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| | 15 - 1667 | 15 - 205 |
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| 9315 922112 9316 920506 9317 916695 9318 463278 9319 881505 9320 920349 9321 955281 9322 719113 9324 727465 9324 727465 9326 692781 | | 15 - 881 | 15 - 645 | 15 - 562 15 - 579 | 15 - 371 | 15 - 627 | 15 - 356 15 - 522 | 15 - 630 15 - 940 | 15 - 462 |
| 9315 9317 9318 9319 9320 9321 9322 9322 9322 9324 9325 | | 1 - 867 | 1 - 631 | 1 - 548 1 - 565 | 1-357 | 1 - 613 1 - 644 | 1 - 342 1 - 508 | 1 - 616 1 - 926 | 1 - 448 |
| | | 922112 | 920506 | 916695 463278 | 881505 | 920349 955281 | 719113 909182 | 727465 959620 | + |
| | | 9315 | 9316 | 9317 | 9319 | 9320 | 9322 | 9324 | 9326 |
| | | HCFMZ90 | HCFMW71 | HCFMU01 HCFMP46 | HCFMO47 | HCFMJ37 | HCFMH46 HCFLY85 | HCFLY53 HCFLY08 | HCFLV30 |

| AA002042, AA001671, AA001672, AA694015, AC009263, and AR068552. | T64674. | AA984238, AA525888, and AA525391. | W31316, AA019148, and AC004706. | AA568314, AA610433, AI090334, AI871691, AA629679, AL044339, AA515677, AA846923, AI799421, AI049701, AI368732, AA523833, AA173334, AL039309, AI751191, T08386. | AW089016, AI821722, F11421, AI732162, AI557422, AW265688, AI049630, AI821172, | AI125167, AA662614, F31811, AA663074, AL046746, AA984829, AI792063, AA706202, | AA586646, AA130647, M78021, AA862029, AA229988, AW272640, AA610255, AA368012, | T48809, AI130709, H96146, AI536858, AW089950, T41134, AI457389, AI537995, AI719142, | AA158549, AI814682, AW190277, AI636734, AA564682, F31867, AI291439, AA524846, | AW023111, AA481760, AA654482, C06028, AI302156, AA020882, AI054339, AA715351, | H24957, AA728861, AA807704, AL079734, AI564301, H67064, AA878492, AI370170. | AA557945, AI802526, AI783911, AI671077, AI114494, AL134332, AW166920, AA856817. | W63553, AW151935, AI160786, AA434078, T17332, AA384064, AA599080, AW188742. | AA323085, H40478, AA371580, R70883, AW419389, H43183, AA368329, AA569089, AJ907046. | AA126635, AI342677, AA598663, AI342786, AI300413, AW148775, AA326330, AA371410. | AA102737, R83585, AW381823, AI376454, AA828463, D51877, H54601, T65895, AI431513. | H58393, C18779, F08866, AW162750, AA595504, AI301373, AI929796, AA702618, AA357878, | AA315361, AC005015, Z93016, AC005081, AP000512, Z83844, AC004458, AC004796, | AC006966, AC004033, AC005189, AC006946, AP000547, D84401, AL035400, Z95116, | AP000558, AP000501, AL139054, AL049776, AC005102, AL031666, AC002477, AC005519, | AL031055, AC003070, AC005180, AL031281, AC005088, AC005529, Z98200, AL008725, | AC003950, AL031311, AL021391, AC005089, AL034429, AC004884, AL136295, AC002301, | AL031432, AC002312, AC007376, AC002300, AJ003147, AC003692, AC004815, AL096701, | 295115, Z68277, AC002425, AF111168, AP000067, AC005702, AL034417, AP000557, | AC005046, AC005500, AL049869, AC004686, AL022476, U91326, AL008582, AC002550, | AP000503, AL031587, AC005412, AC005280, AC005911, AC006251, AL031230, U85195, | AC006241, AL049694, AL049636, AL109839, AL022329, AF134726, AC009516, AL109827, | AC005193, AL031681, AC002565, AC002400, AC007386, AC005049, AL031280, AE000658, | AC004650, AC004905, Z68870, AC004216, AC004791, AC005197, AC005332, AL031228, | AC000353, AF030453, AC006064, AC006511, Z98304, AC007842, AC005520, AC002357, | AC005839, AL050348, AC005094, AC002351, AC005696, AC005800, AF001549, AC005484, | AC008009, AC002039, AC004953, AC005736, Z93930, U96629, AL020993, AC000379, | AC006132, AF047825, AC004832, AP000065, AF196779, AC005209, AL034555, AP000116, | AC003101, AC007546, AC004881, AC007160, AC006538, U82828, AP000553, AF196970, | AP000589, AP000210, AP000132, AC004659, AL035458, U80017, AL121825, AF064861, |
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| 15 - 560 | 15 - 502 | 15 - 837 | 15 - 774 | 15 - 435 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| 1 - 546 | 1 - 488 | 1 - 823 | 1 - 760 | 1 - 421 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| 785560 | 850403 | 850406 | 964792 | 502414 | | | | | | | | | _ | | | | | | | | | _ | | | | | | | , | | | | | | |
| 9327 | 9328 | 9329 | 9330 | 9331 | | | | | | | | | ••. | | | | | | | | | | | | | | _ | | | | • | | | | |
| HCFLT86 | HCFLM03 | HCFLL03 | HCFLK10 | HCFLB24 | | | | | | | | | • | • | | | | | | | | | | | | | | | | | | | | | T |

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| | 15 - 230 | 15 - 542 | 15 - 380 | 15 - 467 | |
| | 1 - 216 | 1 - 528 | 1-366 | 1 - 453 | |
| | 779327 | 850413 | 772256 | 850435 | |
| | 9332 | 9333 | 9334 | 9335 | |
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| A | 15 - 350 A | 1 | | 15 - 701 A | <u> </u> | <u>Z</u> · | <u> </u> | <u> </u> | 15 - 456 T | \vdash | A | TH. | A | ¥ | <u> </u> | | + | - | 1 | | | \dagger | - | 1 | | | 15 - 659 A | 15 - 304 | 15 - 454 R6 | 15-419 RI | 15 - 222 HG |
| | 1-336 | 1 - 342 | 1 - 387 | 1 - 687 | | | | | 1 - 442 | 1-451 | | | | | | | 1 150 | 1 - 452 | 1 - 443 | 1 - 362 | 1 - 463 | 1 - 433 | 1 - 423 | 1 - 400 | 1 - 236 | 1 - 375 | 1 - 645 | 1 - 290 | 1 - 440 | 1 - 405 | 1 - 208 |
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| | 9353 | 9354 | 9355 | 9356 | | | | | 9357 | 9358 | | | | | | | 0360 | 6006 | 9300 | 7301 | 9502 | + | 400% | | \dashv | -+ | \dashv | 9369 | \dashv | _ | 9372 |
| | HBUAG44 | HBUAC23 | HBTAE84 | HBMXU73 | | | | | HBMXS23 | HBMXL78 | | | | | | | UDMAI 41 | UDMV770 | HDMAJ/0 | HDMVE72 | HBMW713 | UDMINIOTE STANKE | TOWNS CO | TIDIM W K02 | HBMWK06 | HBMWI28 | HBMWE35 | HBMVQ26 | HBMV041 | HBMVI33 | HBMVG85 |

| D59551, D59317, D80522, D51103, D59927, AW179332, AW360811, D80024, C14046, AI535686, D80193, D58101, D80302, C05763, AA514186, AA514188, AW178906, AI525912, AW369651, D45260, AI525242, AI557774, AI525235, AI557751, AI525920, AI525917, AA514184, AI535959, AI525215, AI525237, AI525237, AI525923, AI525926, T11191, L81613, A87015, AR008278, A84916, AB002449, A62300, A62298, I14842, AB028859, A82595, AR054175, AR060385, AJ132110, AR008277, AR008281, AR018138, AF058696, X68127, X67155, D88547, AR016691, AR016690, and U46128. | W38346. | | R78567, R22348, AI138749, AI432160, and AI378357. | T89785. | AA252232, H30735, H30819, and AF085888. | AA766345, AA258762, AA258882, and R87385. | | T80409. | AA160470, AA594945, AL047220, and Z98884. | R09103. | AA077388, AL043009, AF039185, AW243793, AA974503, AL079869, AL044339, AA078338. | AL079812, AI567674, AW008212, AI499938, AL039996, AI732120, AA908687, AW102955. | AI679782, AI582769, AL041706, AA563818, AA255853, AW406162, AA287570, AA680243, | AL044340, AA884598, T74524, AI356440, AI110760, AL121235, AA578590, AL038842. | AI799642, AW089016, AA349366, AA806796, AI635279, AI434695, AI917156, AA643434. | AI053790, AW302711, D82290, AI144081, AI313166, AW276586, AI708009, C06151, | AW301906, AW342042, AL044940, AL048925, AW302753, AC004878, AC005081, AC004895, | AC010205, AC004134, AC003682, AC005488, AC002350, AL023553, AC006057, AL022165, | AC008101, AF053356, AC003104, AL080243, AL121653, AC005783, L78810, AC005015, | AL139054, AC008079, AL133448, AC005088, AC005399, AL034423, AP000557, AF001549, | AC007731, AL049869, AC007371, U85195, AF196779, AC004882, AC004814, AC005520, | AL022322, Z99716, AC002563, AL121603, AE000658, AC000025, AL050318, AC004883, | AL022238, AL022721, AL096712, U95740, AC007055, AF134726, AC004796, AC004815, | AC004217, AC006130, AC005940, AC005531, AC006511, AC006014, AC006571, AL096791, | AC007919, AC007227, Z97630, AC004000, AP000350, AL049712, Z84487, AC007066, | AC002404, AF088219, AL050307, AC009247, AC006285, AC002115, AL008582, AC000026, | AC003007, AC004019, AC004832, AL035458, AL135744, AF165926, D87675, AL080242, | AC005500, AC004542, Z98051, AL021154, AC002543, AC002303, AL022323, AP000690, | Z93241, AC005606, AC005067, AC007324, AC004033, U80017, AC004638, AF111168, | ALU49//6, ACU06006, AP000045, ACU06111, ALU22311, AC008040, AP000338, AC006251, |
|---|----------|----------|---|----------|---|---|----------|----------|---|----------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| | 15 - 530 | 15 - 352 | 15 - 667 | 15 - 439 | 15-611 | 15 - 1002 | 15 - 373 | 15 - 421 | 15 - 609 | 15 - 373 | 15 - 901 | | | | | | | | | | | | | | | | | | | |
| | 1-516 | 1 - 338 | 1 - 653 | 1 - 425 | 1 - 597 | 1 - 988 | 1 - 359 | 1 - 407 | 1 - 595 | 1 - 359 | 1 - 887 | | | | _ | | | | | | | | | | | | | | | |
| | 692733 | 677209 | 625627 | 792736 | 731802 | 678164 | 888393 | 783043 | 646279 | 731800 | 911168 | | | | | | | | | | | | | | | | - | | | |
| | 9399 | 9400 | 9401 | 9402 | 9403 | 9404 | 9405 | 9406 | 9407 | 9408 | 9409 | | | | | | | | | | | | | | | | | | | - |
| | HBMCZ30 | HBMCT24 | HBMCT09 | HBMCS92 | HBMCS55 | HBMCR25 | HBMCQ95 | HBMCG84 | HBMCF20 | HBMCB55 | HBMBX72 | | | | | | | | | | | | | | | | | | | |

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|---|-------------------------------|---|-----------------------|---|---------------------|---|
| | 15-613 | 15 - 674 | 15 - 555 | 15 - 297 | 15 - 534 | 15 - 952 |
| | 1 - 599 | 1 - 660 | 1 - 541 | 1 - 283 | 1 - 520 | 1 - 938 |
| | 746633 | 668981 | 973805 | 783520 | 723094 | 912789 |
| | 9410 | 9411 | 9412 | 9413 | 9414 | 9415 |
| | HBMBX64 | HBMBX55 | HBMBU52 | HBMBN85 | HBMBI49 | HBMBF89 |

| | | | | | T86779, W72654, T86785, AI356251, AW134759, AA761197, T32285, R52494, C00788, |
|----------|------|--------|---------|----------|---|
| 3 | | | | | AA983316, AI/97525, AW135739, AA213524, W49846, AA938126, W76616, AI419595, T86686, AA635708, AI962906, T20097, T30685, H59977, AA565271, AI088829, AW418557, U83525, AA481982, AA482087, AA864361, and Z22819. |
| HBMBF78 | 9416 | 623421 | 1 - 950 | 15 - 964 | AA683395, W39500, and AA159553. |
| \dashv | 9417 | 879588 | 1 - 388 | 15 - 402 | AA770612. |
| \dashv | 9418 | 859500 | 1 - 339 | 15 - 353 | W73386, and AI910369. |
| HBMBB57 | 9419 | 734915 | 1 - 324 | 15-338 | AW080136, AA648683, AI625125, AW193143, AA635976, AI064811, AA935848, AL037910, |
| | | | | | AA564865, AA627154, AW166653, AI932443, AA501614, AW302709, AA578231, AA491807. |
| | | | | | AA845804, AI565097, N68677, AI572115, AI302917, AA857381, AI560195, AL041375, |
| | | | | | AL040054, AA533344, AA515046, AA366716, AL047429, AI769271, AA229935, AI024339, |
| | | | | | AW028908, AI287706, AA757888, AI539530, AA508873, AI061313, AW020094, AA833896, |
| | | | | | AW117860, AA833875, AI249688, AC006023, Z84469, AC004019, AC005412, AL121603, |
| | | _ | | | AC006487, AP000509, AC004217, AP000512, Z98950, AP000359, AC008101, AC006077, |
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| | | | | | AF003626, AP000050, AF067844, AC016025, AC006008, AL022476, AC005215, AF111169, |
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| | | | | | Z84487, AL031427, Z84466, AC004675, AC005914, AC006449, AC007226, AP000511, |
| | _ | | | | AC005399, AL031985, AC007934, AC006115, AL050308, AC005839, AF196969, U95742, |
| | | | | | AL023553, AF109907, AC004914, AC006111, AC002105, AL008725, AC005058, AC005191, |
| | | | | | AC003950, AL022320, AC003029, AC004686, AC004843, AL133163, Z93023, AP000047, |
| | | | | | AC00/216, AC002430, Z83822, AP000432, AL034421, AC007384, AC004834, AC006239, |
| | | | | | AL034420, AC00/664, AL121825, AC006312, AL031846, AF064861, Z86090, AC004875, |
| | | | | | AL008/13, AL049339, AC0046/8, AC000032, Z83986, AL034423, AC006972, AC005726, |
| _ | | | | | AC00303/, AC003944, AL008582, AC006146, AC006384, AC004531, AC003690, AF121781, |
| | • | | | | AC005180, AC007649, AP000692, AC006511, Z94793, AL049694, U95743, AL139054, |
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| | | | | | AP000115, AC004216, AC005632, AL035249, AC002551, AC006501, AC004000, Z82181, |
| | | | | | AF107885, AC006966, AC005088, AL009179, AF088219, AC006530, Z84480, AC005618, |
| | | | | | ACUUSU/9, ACUUSS32, ACUUS800, U85195, AC002040, AF190465, AP000502, AC004996, |

| AC006318, AL033376, AC005694, AP001052, AC005696, D84394, AC005086, AL022165, AC004841, AC004647, AL049780, AC005207, AC007050, AC007041, AC003664, AE000658, AC004929, AC003663, Z98884, AL024509, AC004967, AC005740, D87675, D28126, U91326, AL023280, AC006084, and AC001231. | AJ242976, | AF037641, and AA321292. | AA737327, and AI433448. | R93012, and AA856773. | AA460141, AA460739, AA487420, AA487676, AI620475, AW014765, AA826244, AI807477. | AI859817, AI888070, C01748, and AA487462. | H19011, AA017590, H86398, AA969667, AA055983, and AL009182. | H10282. | N55533. | H72483, AI125256, and C15483. | AI566099, AW027347, AI921989, AA769157, AA722358, N66761, AI696509, AI825693. | AA651815, AI933564, D59982, D61092, AA860600, D59981, D61152, AI873970, AA831741, | H02809, AA653665, and AL050064. | W17090, and AW022657. | | N80651, and AC002400. | H37828, and AA707167. | T56980, AF178030, and AF147313. | AA290916, R00373, AW387125, AA449392, AW387122. and AL080141. | H38564, AC005049, AC005245, AC006014, AC005488, AC006195, AC003043, AF001549, and AC005071. | AA084033, AA126948, AA126949, AA070285, U67770, X02414, X63389, and AA070164 | C14331, C14429, D59275, C14389, AA305578, D80227, D59859, D80164, D80378, D80195 | D59467, D59502, D80038, D80269, D51423, D58283, D80022, D51799, D80166, D80253, | D59619, C15076, D80210, D80391, D80240, D80043, D59787, D81030, D81026, D80024, | D80212, D50979, D80193, D80196, D80188, AA305409, D80219, AA514188, D57483, D59927, | D80366, D51022, D59610, D59889, D50995, D80045, D80248, AW177440, AA514186, D80241, | AW360811, AW178893, D80522, D80251, D51060, C14014, D80133, T03269, AW375405, | AW1/9328, C/5259, AW1/9020, AA809122, C14407, AW179023, AW366296, AW378532, | AW178015 AW177501 AW177511 COSSOS DOODD AW170010 AW17016 AW17551. | D51250, AW352171, D80439, AW377676, AW178906, AW352170, AW177731, D80132. |
|---|-----------|-------------------------|-------------------------|-----------------------|---|---|---|----------|----------|-------------------------------|---|---|---------------------------------|-----------------------|----------|-----------------------|-----------------------|---------------------------------|---|---|--|--|---|---|---|---|---|---|---|---|
| | 15 - 357 | 15 - 435 | 15 - 366 | 15 - 498 | 15 - 819 | | 15 - 965 | 15 - 472 | 15 - 385 | 15 - 636 | 15 - 447 | | | 15 - 465 | 15 - 471 | 15 - 304 | 15 - 439 | 15 - 442 | 15 - 412 | 15 - 440 | 15 - 430 | 15 - 1454 | | - | | | | • | <u> </u> | |
| | 1 - 343 | 1 - 421 | 1 - 352 | 1 - 484 | 1 - 805 | | 1 - 951 | 1 - 458 | 1 - 371 | 1 - 622 | 1 - 433 | | | 1 - 451 | 1 - 457 | 1 - 290 | 1 - 425 | 1 - 428 | 1 - 398 | 1 - 426 | 1 - 416 | 1 - 1440 | | | | | | - | - | |
| | 578817 | 930481 | 915970 | 726475 | 797604 | | 95136 | 727987 | 710880 | 738615 | 787320 | | | - | | 754875 | 777456 | 625319 | 765352 | 847851 | 849633 | 688006 | | | | | _ | | | |
| | 9420 | 9421 | 9422 | 9423 | 9424 | | 9425 | 9426 | 9427 | 9428 | 9429 | | | 9430 | 9431 | 9432 | 9433 | 9434 | 9435 | 9436 | 9437 | 9438 | | | | | | | | [|
| | HBMBB45 | HBMAC54 | HBJNC01 | HBJNB52 | HBJMW95 | | HBJMW03 | HBJMU53 | HBJMS40 | HBJMR59 | HBJML90 | | | HBJML80 | HBJME11 | HBJMC69 | HBJMA80 | HBJLX09 | HBJLW74 | HBJLU36 | HBJLR37 | HBJLR31 | | | | | | | | |

| AW178907, AW178775, AW179024, D80247, AW179018, AW177505, D80134, D58253, D51103, AW352158, AW360841, AW178909, D59373, AW352117, AW177456, AW179329, AW176467, AW369651, AW178980, AW177733, AW378528, AW178908, AW178754, AI555686, AI910186, AW352174, AW177904, D59695, D52291, AW1779012, AW178914, AI535686, AI910186, AW352174, AW177904, D80157, AW177722, AW177723, T11417, D80168, AW367967, AW179009, D51759, C06015, AW1787721, AW1778781, AW360834, D58246, T4827, C14344, D81111, C14298, AW1787911, AW378543, AI905856, AW352163, C14227, C14344, D81111, C14298, AW177723, AW378784, AW378549, AW378539, AW378540, AW177723, AW378540, D59653, D59653, D59673, D89673, D896773, D89673, D89677, D89674175, AR066488, D88677, D89674175, AR066488, D88677, D8967477, AR066488, D88677, D8967477, D8967477, D8967477, D89678, D89679, D89678, D89678, D89678, D89678, D89678, D89678, D89678, D89679, D89678, D89679, D89678, D89678, D89678, D89678, D89678, D89679, D89678, D89678, D89679, D89679, D89679, D89679, D89679, D89678, D89679, D89678, D89679, D89679, D89679, D89678, D89679, D8 | AA192415, A1216292, AW151132, A4573503, AA8035063, AKU32003, and AKU08382. | AI720923, AI628097, AI149416, AI432468, AI038395, AI360839, AA235040, AA010019, AW104170, AI266742, R80513, AA954623, R80407, H81294, AA235207, AA992698, T98403, AA480593, AI352046, AA807246, AA010161, T98402, AW078986, AI572359, R64407, AI547277. and AC004148. | AI763332, AI744915, W02174, W32586, and AI.109677 | AI739538, AC006140, AB023052, and AP000513. | AA278175. | | AW162288, AI696793, F13749, AI431434, AI537538, AI358089, AW021116, AW157005, | AA657835, AA508809, AW328331, AA130647, AI954525, AA992126, AI051037, AI028510, AI751162. AW271917, AI340641, AA728937, AI500552, AI821881, AI821018, AA665252 | AA629412, AA829065, AA804925, AI566408, H90844, N67343, AW089589, N23260, H71678. | D29500, AL046746, H79308, AL038936, F30251, AI291961, AI887716, AW192065, AI571656, AI309384 A1961536 AA714011 A1779417 AA644320 AI508060 AA66600 AA66 |
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| | 15 - 337 | 15 - 1007 | 15 - 697 | 15 - 820 | 15 - 399 | 15 - 407 | 15 - 355 | | | |
| | 1-323 | 1 - 993 | 1 - 683 | 1 - 806 | 1 - 385 | 1 - 393 | 1 - 341 | | | |
| | 614915 | 847853 | 772000 | 952843 | 919446 | 771975 | 746398 | <u> </u> | | |
| | 9439 | 9440 | 9441 | 9442 | 9443 | 9444 | 9445 | | | |
| | HBJLF04 | HBJLE70 | HBJKI77 | HBJKG07 | HBJKF02 | HBJKD77 | HBJKD64 | | | |

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| AW307050 AI570043 AA563770 AA271512 ACMOSA12 AT117526 ACMOMAC 70000 | AL133245, AC004692, AL022313, AL050332, AC000353, AC006236, AC005280, AL031255, | AC005726, AC005899, AC004832, AC020663, AL021939, AL035413, AC000052, AC005539, | AC004322, AC00544, AC004013, Z82244, AF000270, AC004854, AC005884, AC005996, AP000032, AC006080, AC005874, AF134471, AC005081, AL021453, AP000215, AP000116 | AC007057, AL049872, AC006356, AL049757, AC005231, AC007055, M87889, AL035587, | AC004167, AC005696, AC005531, U53930, AP000557, AP000558, AC009516, AC006468, | AL022238, AC004996, AC005399, AF038458, AL031296, AC006600, AC003667, AC007283, | AC005015, U91321, AC004072, AC004534, AC007899, AL021154, AL035667, AL031003, | AC007308, D87675, AP000513, AC005800, AC002483, AC004794, AC006064, Z68321, | AC005013, L35485, AC008071, AC005962, AC004921, AL121653, AL049780, AC002115, | ACUUUU/U, AFUUUSU3, ACUUSU3/, ACUU/226, ALU3337/6, ALU335448, ACU02375, AC004087, | AC004963, AC005900, AC005520, AC005670, AC007376, AC00548, AC005080, AC005403 | AC002350, AL080285, AC007237, AL031407, Z99716, AC009509, AL034493, AP000441, | AL031393, AC004887, AL050318, Z83844, U85196, AC004452, AC005616, AL050341. | AP000311, AL008719, AF011889, AF111169, AF196779, Z72521, AC006530, AL034553, | AC006512, AC005193, AC005057, AC004150, AL031733, AC005062, AC006277, Z95114, | AC005529, AC007731, AC005602, AC006057, AF050154, Z84476, AP000689, AC003682, | AC002059, AL008735, AP000090, AC004477, AC006111, AC002530, AP000356, AL133485, | AC003665, AE000661, AC007228, AC004890, AL109984, AL020997, AC004253, AC004859, | AL022165, AL022163, AC018633, AF195658, AC005192, AC006071, AC004876, AF031078, | AL049830, AC005914, AC004263, AL034555, AF030876, AC004687, AC005004, AB003151, | ACU05229, ACU05664, ACU05500, AL021395, AC000025, AL034402, AC003690, AC005876, | AL022397, ACU00097, ACU05780, AF109907, AC005527, AC006547, AC000041, AC002091, | Z467/3, AJ003147, AC005209, AC002425, AL035089, AL031589, AC004895, AL035681, | ALU31053, Z8Z1/3, AC004408, AC006270, AC005212, AF111167, AC006132, AC003955, | R77177 | AA303007, AA630672, AA634147, N22465, AI249447, AA679532, AA478048, AI821670. | AI357823, AA613249, AW274349, AI524360, AI254913, AI696962, N22023, AI590906, | AW264934, AA012982, AA688359, AI791913, AW270258, AI792133, AW303196, AA169245. | AA580808, AI821714, AA528554, AA631517, AL119552, AA188664, AW301350, AA631507, | AA483771, AW304580, AA480574, AA112924, AL048135, H73863, AI580250, AW270270, | AC006455, AL121754, AP000350, AC002425, AC004778, Z83844, AP000155, AF111168, | AC002540, AF190465, AL109/98, AC007312, AL121655, AC004859, AC006077, AC002369, | AC0048/6, AC002126, AP000151, AP000109, AC007686, AP000501, AC002472, AP000690, |
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| IT SOURCE IT DOSESSES IN TAXABLE SECTION OF THE SEC | AL109884, ALU35587, AL078602, AL031733, AL009183, AL031311, AC006040, AC003663, AC007664, AC000353, AF165175, AC005632, U91323, AC006088, AP000113, AP000045, AC002303, AF165926, AF117829, AC005322, U91323, AC006082, AC004790, AF109907, AC006121, AL078584, AF207550, AP000039, AF068006, AL121658, AC004895, AC006241, AF184110, AC005015, AL034421, AC002477, AC005102, AC004895, AC004881, AC007285, AC005666, AC005037, AC002407, AC002407, AC005027, AC005003, AF1031659, AC006001, AL022238, AC005565, AL021154, U52112, AC006411, AL031659, AL021326, AC006511, AC006001, AL121653, AC005607, AC006407, Z82244, Z92547, AL080243, AC006406, AL121653, AC006406, AL033527, AL033527, AL034976, AC006409, AC006599, AC006409, AC006409, AC006409, AC006599, AC006409, AC006599, AC006599, AC006509, AC006 | , and 201100, and 201100, and 201100, | | AA256446, AI806202, AA746104, and AA811237. | AI445151. | AA477047, AW383764, AA321993, AA309810, W25768, AA465618, R24995, W07130, R50327, AA368102, N58603, AA356448, AW015964, W69552, AI791923, AA525463, AA526678, AI792408, AC005071, AF161081, and AF161080 | AA113985, AA010096, AA333937, and UZ6150 | Z78294, AA808842, AA715173, AA715075, AC005102, AC006449, AC005697, AL031311, AC007277, AC004686, AC009247, AC006211, AL096701, AL080242, AC004125, AC005914, AF207550, AC004659, AC005746, AC004491, AC006947, AL023879, AC006487, AL020997, AC005940, AC004827, AC006115, AC016025, AC005189, AF196972, AF109907, and | AC007057. AA285172. |
| | | 15 - 427 | 15 - 365 | 15 - 566 | 15 - 542 | 15 - 393 | 15 - 618 | 15 - 569 | 15-215 |
| | | 1-413 | 1 - 351 | 1 - 552 | 1 - 528 | 1 - 379 | 1 - 604 | 1 - 555 | 1 - 201 |
| | | 847857 | 710931 | 771982 | 916077 | 835733 | 894345 | 919371 | 937583 |
| | | 9448 | 9449 | 9450 | 9451 | 9452 | 9453 | 9454 | 9455 |
| | | HBJKD49 | HBJKD40 | HBJKA77 | HBJKAUI | HBJJX53 | HBJJT12 | HBJJN02 | HBJJL10 |
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| 1 4 11737/4108 | AI820773, H30102, H39985, H21775, AI668592, AI821454, R40787, H38487, and ACROSTAGE | AI088489, AA194062, AI123551, AA759303, AW028108, AI093432, AA836264, AA767044, AI274044, AI434773, AW135037, AI808043, AA806714, AW087576, AA883729, AA809688, AA84703, AI524547, AA865016, and A Condosa | 104202, | AI091292, AI005391, AA806165, AI002979, AI702990, AA158537, N54188, AA069178, AA091807, AA095638, and AC006146 | | | | | AA147107, AA425502, R79257, H03146, W58353, C19029, AA657402, AI050680, AA757308, AI063898, AA95368, AA360708, AI303330, III063898, AA953682, AA360708, AI303330, III063898, AA963898, AA963898, AA963898, AI063898, AI06388, AI0688, | 41148152 D51354 D80752 D80042 AWA4567 D50257 D80316 D50216 | D80210. D51423. D80134. D50619. D801913. AM 242391, D39101, D80219, D39213, D80221, D80240, | 039156 AI 043441 AI 030150 AI 030031 | T24119 AI 039678 AI 039074 AI 03827 | T24112, AL037726, AL038531, AL039566 | AL039109, AL040992, AL039924, AL039128, AL044407, AL039659, AL036973, AL03986 | AL045337, AL037051, AL045353, AL036725, AL039423, AL045794, AL045341, AL039410 | AL042909, AI535983, H00069, AL038025, AL044530, AL043422, T11051, D81026, AI 043423 | D80045, D50995, C14014, C75259, R47228, AL036418, AI535783, AW013814, AW452756 | D59889, AW451070, C15076, AL037526, AL037639, D80022, D80195, AL036196, D80038 | AL037615, AL038851, AL037082, T23659, D81030, D58283, T11417, T02921, D80188, D51799 | D80378, D59467, AL036924, F13647, AL036117, AI557751, AL036767, T03269, AL036679. | AL037601, AL036238, D80522, D80212, D50979, AL036190, AL036733, C14298, T48598, | AL036964, Z21582, D59502, AA285331, C14429, D80164, AA514190, D59695, AL036158, | D80166, AL037027, D59859, D80269, D80268, AL037054, AL037178, D52291, Z25782, D58253 | AL036191, D80024, AL036765, AL036227, D57483, Z99396, D59610, C14389, D59627, D80241 | D81111, AL036998, AI910186, AL037177, C14331, AL036207, D51060, H00072, AA305409 | AL037021, AW178893, AL036174, D51079, AL037047, AW450376, AW177440, AL036139 | D51022, AW179328, AL036167, D80014, AW178775, AA305578, AW378532, AL036132. | D51213, AW352158, AW377671, AI905856, AW369651, D80251, D80064, D51097, AA514188. | D80248, AW178762, AL037077, AW177501, AW177511, AL037002, AW360834, AI557774. | AA514186, D80133, AW360811, T02974, D80302, AW352117, C05695, C14407, AW176467, AW375405, AW378540, 725783, AW366206, D80133, AW366306, DR0133, AW366506, DR0133, AW366506, DR0133, AW3666006, DR0133, AW3666006, DR0133, AW3666006, DR0133, AW3666006, DR0136, DR013600, DR0136, DR013600 |
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| AT142126 R5454 H24334 AT142007 224 A 17274108 | AI820773, H30102, H39985, H21775, A1668 | AI088489, AA194062, AI123551, AA759303, AWC AI274044, AI434773, AW135037, AI808043, AA80 AA847003, AI524547, AA865016, and AC004082 | R35568. | AI091292, AI005391, AA806165, AI002979 AA091807, AA095638 and AC006146 | AA729724. and AC006116 | H69862. | T79551, T86369, R98540, and T98454 | H69017, and H69779. | AA147107, AA425502, R79257, H03146, W58353, C19029, AA657402, A105066 | A1148152 D51250 D80252 D80042 A4124 | D80210, D51423, D80134, D50419, AW24 | T23947, D59927, AL039538, AL039564, AL039566, AL043441, AL043441, D8 | AL043445, AL039509, D80366, AL039108 | AL039625, AL039648, AL039629, D80168, T24112, AL037726, AL038531, AL039566 | AL039109, AL040992, AL039924, AL03912 | AL045337, AL037051, AL045353, AL03672. | AL042909, AI535983, H00069, AL038025, A | D80045, D50995, C14014, C75259, R47228, | D59889, AW451070, C15076, AL037526, AI | AL037615, AL038851, AL037082, T23659, I | D80378, D59467, AL036924, F13647, AL036 | AL037601, AL036238, D80522, D80212, D50 | AL036964, Z21582, D59502, AA285331, C14 | D80166, AL037027, D59859, D80269, D8026 | AL036191, D80024, AL036765, AL036227, I | D81111, AL036998, AI910186, AL037177, C | AL037021, AW178893, AL036174, D51079, | D51022, AW179328, AL036167, D80014, AV | D51213, AW352158, AW377671, AI905856, | D80248, AW178762, AL037077, AW177501, | AA514186, D80133, AW360811, T02974, D8 AW375405 AW378540 725783 AW366206 |
| 15 - 578 | 15 - 709 | 15 - 1056 | 15-413 | 15 - 790 | 15-531 | 15 - 463 | 15 - 687 | 15 - 545 | 15 - 639 | 15 - 1059 | | | | | | | | | | | | | | | | | | | | | |
| 1 - 564 | 1 - 695 | 1 - 1042 | 1 - 399 | 1 - 776 | 1-517 | 1 - 449 | 1 - 673 | 1 - 531 | 1 - 625 | 1 - 1045 | | | | | | | | | | | | | | | | | | | | | |
| 716409 | 699059 | 916078 | 597100 | 861363 | 930448 | 725081 | 697753 | 754891 | 784267 | 971377 | | | | | | | | | | | | | | <u> </u> | | | | | | | |
| 9456 | 9457 | 9458 | 9459 | 9460 | 9461 | 9462 | 9463 | 9464 | 9465 | 9466 | | | | | | | | - | | | | | | • | | | | | | | |
| HBJJE44 | HBJJA32 | HBJJA01 | HBJIL42 | HBJIL36 | нвлн60 | HBJIF51 | HBJIF31 | HBJIB69 | HBJIA80 | HBJHX73 | | | | | | | | | | | | | | | | | | | | | |
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| AW37834, AW17932, AW37675, AW179023, AW17890, BR0247, AW17731, D59273, AW37834, AW17929, AW377676, AW372171, AW17890, AW352170, AW17731, D59373, AW360841, AW17020, C06015, AW179024, AW177201, AW17731, D59373, AW360841, AW17020, C06015, AW17902, AW177246, T03116, AW177322, AW17890, AW178980, AW17890, AW1791746, T03116, AW172922, AW17890, AW178980, AW17891, AW17733, AW37828, A25909, A67220, AR022207, X68127, A83396, A88477, A86792, D3646, LW1820, AW17731, AS6903, A67220, AR022207, X68127, A83396, A884773, A88772, D3646, AW1819274, AR067731, AR067821, AR067821, AR067821, AR067821, AR067821, AR067821, AR067821, AR067821, AR06782, A88222, AB120969, X73004, A88173, AR068206, AR018924, A63067, A51047, A63064, AR018923, A48774, A63072, A48775, AR068206, AR018924, A63067, A51047, A63064, AR018922, A48774, A63072, A48775, AR068206, AR018924, A63067, A51047, A63064, AR018922, A48774, A63072, AR06940, AZ04006, AZ04005, 119516, A18053, 106839, A3384, A75888, AR064801, A26491, A26401, A26491, A264 | HBJHW74 9467 765380 1-435 15-449 AA716676, H63334, H60316, AI744524, AA836358, Z95116, AC005529, U91321, AC004832, | 1 - 544 15 - 558 | 9470 828996 1 - 424 15 - 438 | (C) 1000 111 (C) 111 (|
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|---------|------|----------|---------|----------|--|
| | | | | | 154669, AI814941, AA680075, AA852300, AA131298, AI686560, AA600068, AI380408, AI694164, AA338172, AA853557, AA304609, AA332521, R94291, AA131543, AW198197, W68749, N56676, AA377810, H58556, AA055184, W58726, AA570322, AA347235, R75738, AA419374, W51908, W52879, A1780505, A A 256330, A 1577400, A 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 |
| HBJHG02 | 9471 | .919450 | 1-643 | 15 - 657 | AA126455, and AA805265. |
| HBJGT03 | 9472 | 923800 | 1 - 497 | 15 - 511 | AW014944, AI571386, AI494432, AW149677, and AC007682. |
| HBJFR45 | 9473 | 722105 | 1 - 519 | 15 - 533 | AW205972, and R06326. |
| HBJFQ08 | 9474 | 959190 | 1 - 326 | 15 - 340 | |
| HBJFP52 | 9475 | | 1 - 471 | 15 - 485 | R33062, and R26486. |
| HBJFP08 | 9476 | - | 1 - 474 | 15 - 488 | AA463369, |
| HBJFK85 | 9477 | 783432 | 1 - 292 | 15 - 306 | AL036438, and R43453. |
| HBJFI48 | 9478 | \dashv | 1 - 489 | 15 - 503 | AC005180. |
| HBJFI30 | 9479 | | 1 - 440 | 15 - 454 | H83540, H83349, T98751, and T89066. |
| нвлен60 | 9480 | 740543 | 1 - 552 | 15 - 566 | AI631029, N51923, H64804, AW016151, N51665, N53749, AA503053, AA917715, R81860, and |
| 4 | | | | | N50398. |
| HBJFH06 | 9481 | 935587 | 1 - 675 | 15 - 689 | AA189094. |
| | 9482 | 711591 | 1 - 638 | 15 - 652 | AA120846, N94144, and AA120845. |
| HBJFD21 | 9483 | 671205 | 1 - 869 | 15 - 883 | W89209, AA236825, T90638, AA255865, AW137746, AA814395, AA814416, AA825532, |
| | | | | | T83165, and T82244. |
| HBJFC23 | 9484 | 423885 | 1 - 616 | 15 - 630 | AW276648, R21362, H10771, AW376712, AI907952, AW407979, AW366716, H08663, and |
| HBJFB10 | 9485 | 968162 | 1 - 421 | 15 - 435 | AI040621, H71636 TR0789 and AI 035086 |
| HBJEY89 | 9486 | 787044 | 1 - 309 | 15 - 323 | W59951. |
| HBJES09 | 9487 | 487020 | 1 - 547 | 15 - 561 | |
| HBJEP70 | 9488 | 757374 | 1 - 476 | 15 - 490 | R64066, and R64081. |
| HBJEN68 | 9489 | 752251 | 1 - 594 | 15 - 608 | H48332, H53781, and H48241. |
| HBJEM45 | 9490 | 722107 | 1 - 306 | 15 - 320 | N35747. |
| HBJEL04 | 9491 | 615515 | 1 - 480 | 15 - 494 | H90849, H78133, and AI023482. |
| HBJEJ06 | 9492 | 935967 | 1 - 385 | 15 - 399 | H37882. |
| HBJEE72 | 9493 | 766187 | 1 - 516 | 15 - 530 | AI739168, AI243692, R61551, AI423731, R53934, AA426249, T06305, AI423751, AA805323, and AA736849. |
| HBJED51 | 9494 | 485130 | 1 - 485 | 15 - 499 | |
| HBJED05 | 9495 | 935058 | 1 - 459 | 15-473 | T19837, Z42292, W56491, Z43884, R10274, H61521, AA047587, AI557059, AA001460, R21700, R56331 H58106 W68068 A1870218 A1007042 AW301022 H71164 E06067 |
| | | | | | 100551, 1255150, 1100000, 12101, 12101, 12101, 1201032, 11, 1101, 120831, and 2,28831. |

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| | | | W85789, AA693974, AI032087. and W85723 | | | AI018729, AI017735, T94096, AA788760, AC004878, AC006014, AC005488, AC005071, | R96956 | N92225 | H02200. | R92654, R92648, and N59587. | D58283, D80195, D80227, D80391, D80166, D59859, D80043, D59502, D51423, D59619 | D80210, D51799, D80240, D80253, D80196, D80269, D80038, D80188, D8011, D50275 | D80193, D57483, D59927, D80219, D81030, D80022, D50979, D59889, D5610, D80366, | D80378, D80045, D59787, C14429, D50995, D80164, D81026, D80041, D80024, T02260 | C15076, C14389, C75259, C14014, C14331, D59467, D51060, AW178893, AA305409, D80134 | F13647, D51250, D80268, D58253, D80949, AA305578, D80168, C14227, D51079, D81111 | D51022, AW177440, AW179328, D80522, D52291, AW178775, AW378532, D50695 | AW352158, AW377671, Z21582, AI910186, AW369651, AI905856, D80251, AA514188, D80248 | AW178762, AA514186, D51097, AA285331, C05695, C14298, AW177501, AW177511, D80064 | D80133, AW360811, C14407, AW352117, AW176467, AW378540, AW375405, AW360834 | D80132, AW366296, AW360844, AW360817, AW375406, AW378534, AW179332, AW377672 | AW179023, AW178905, AW179220, D80302, AI557751, AW352171, D80439, AW377676. | AW178906, AW352170, AW17731, AW178907, AW179019, AW179024, D59373, D80247. | D80014, AW177505, AW179020, AW360841, AW178909, AW177456, AW179329, AW178980. | AW177733, AW378528, AW178908, AW178754, AW179018, AW352174, T11417, D51103. | 103116, AW179004, AW179012, AW178914, AW378525, D80157, AW367967, AW177722, | AW177728, T02974, D51759, AW179009, AW178774, AW178911, AW378543, AW352163. | D58246, C06015, D59503, D80258, AW178983, AW352120, AW178781, T48593, D51213. | AW378539, D59627, AI557774, D58101, AI535850, AA809122, AW177723, D59653. | AW177508, D45260, C14975, AI525923, AW378533, AW367950, H67866, D59317, H67854 | C03092, D45273, AW177497, AI535686, AW178986, N66429, AW177734, AI525917, C14973 | C14344, D59474, AI525227, D51221, D59551, D60010, AA033512, AI525920, AA514184 | D60214, C14957, C14046, C16955, T03048, AI525242, AI525235, AI525912, AI525925. | AI535961, AW378542, AI525215, C05763, Z33452, AI525237, AI525222, D31458, AJ132110. | A62300, A84916, A62298, X67155, A67220, D89785, A78862, D26022, A25909, Y17188. |
| 15 - 438 | 15-119 | 15 - 521 | 15-612 | 15 - 110 | 15 - 269 | 15 - 397 | 15 - 429 | 15 - 436 | 15 - 400 | 15 - 638 | 15 - 821 | | | | | | | | | | | | | | | | | | | | | | | | |
| 1 - 424 | 1 - 105 | 1 - 507 | 1 - 598 | 1 - 96 | 1 - 255 | 1 - 383 | 1-415 | 1 - 422 | 1 - 386 | 1 - 624 | 1 - 807 | | | | | | | | | | | | | | | | | | | | | | | | |
| 573809 | 573834 | 625943 | 577103 | 839964 | 419107 | 847928 | 677407 | 784654 | 757435 | 678272 | 900955 | | | | · | | | | | | | _ | | | | | | | | | | | | | |
| 9496 | 9497 | 9498 | 9499 | 9500 | 9501 | 9502 | 9503 | 9504 | 9505 | 9206 | 9507 | | | | | | | | | | | | | | | | | | | | | | | | |
| HBJEC67 | HBJEA73 | HBJEA54 | HBJEA38 | HBJEA37 | HBJEA13 | HBJDZ49 | HBJDR24 | нвлоо | HBJDM70 | HBJDL24 | HBJCV85 | • | | | | | | | | | | | - | | | | | | | | | | | | |

| REGISTRED REGI | | Т — — — — — — — — — — — — — — — — — — — | _ | | | _ | | _ | _ | | | _ | _ | _ | _ | | | | | |
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| 9508 876430 1-1780 9509 779645 1-467 9510 954090 1-180 9511 781411 1-617 9512 847952 1-432 9513 796659 1-263 9514 732436 1-414 9515 723164 1-427 9516 724619 1-490 9517 682941 1-239 9518 699695 1-395 9520 722195 1-395 9521 847966 1-486 9522 660552 1-395 9523 720008 1-572 9524 864344 1-534 9526 96357 1-564 | AR018138, D34614, D88547, AR025207, X82626, AB028859, AF058696, AR008278, I82448, X68127, AB012117, Y12724, A85396, AR066482, A44171, A85477, I19525, A86792, U87250, X93549, A82595, AR060385, A94995, AB002449, AR016808, AR008443, I50126, I50132, I50128, I50133, AF135125, AR066488, AR016514, AR060138, A45456, A26615, AR052274, Y09669, A43192, A43190, AR038669, AR066490, AR066487, I14842, A30438, I18367, D88507, AR054175, D50010, Y17187, AR064240, AB033111, A63261, AR008277, AR008281, AR062872, A70867, AR016691, AR016690, U46128, I79511, D13509, A64136, A68321, AR060133, U87247, Z32749, AB023656, U79457, AF123263, Z82022, AR032065, AR060382, X93535, and AR008382. | AI828112, AI762456, AI669527, AW270041, AW005454, AI742020, AI813489, AW055057, AW438593, AA702328, AI379307, AA262137, AA643040, AA694381, AA281632, AW243902, AI422791, AA493261, AW117912, AI034474, AW236597, AW440103, AI292303, AA281692, D19612, AA768827, N99706, AI081788, AI868989, AA514538, AA877024, N71978, R24110, AW449526, R17044, AW082476, AA369546, R24059, AI290151, R26499, AI383270, AA612940, AW082653, R50985, C16425, R26471, AI701667, D62032, AA468881, AA522467, AI371310, N93082, and W24736. | | R02678. | T88832. | AW206286, AW291675, AW025778, T99524, AW026757, and AC005682. | N89814, AA209448, AW239298, AP000541, and AJ239321. | T81741, AA778192, W85729, and AL132777. | N71474. | AI089358, AA056686, AI089564, AI086521, AI669927, AI817821, AA056582, AA847148, AI827763, and AB002370. | | W88646, AI368895, and Z39215. | H53026. | AI479236, AI935423, AA255972, and AA243356. | 228561. | AA429379. | AA252703, AI760942, and AA565583. | H04563, R80434, AA531428, Z82215, and Z95114. | AI810593, and T89570. | AA701314, and AA004793. |
| 9508 876430 9509 779645 9510 954090 9511 781411 9512 847952 9513 796659 9514 732436 9514 732436 9515 723164 9516 724619 9516 724619 9517 682941 9518 699695 9517 682941 9518 699695 9520 660552 9521 847966 9521 847966 9521 847966 | | 15 - 1794 | 15 - 481 | 15 - 194 | 15 - 631 | 15 - 446 | 15 - 277 | 15 - 428 | 15 - 441 | 15 - 504 | 15 - 253 | 15 - 409 | 15 - 505 | 15 - 888 | 15 - 500 | 15 - 409 | 15 - 586 | 15 - 548 | 15 - 389 | 15 - 578 |
| 9508 9509 9510 9511 9513 9514 9514 9515 9518 9518 9520 9520 9523 9523 9523 | | 1 - 1780 | 1 - 467 | 1 - 180 | 1 - 617 | 1 - 432 | 1 - 263 | 1 - 414 | 1 - 427 | 1 - 490 | 1 - 239 | 1 - 395 | 1 - 491 | 1 - 874 | 1 - 486 | 1 - 395 | 1 - 572 | 1 - 534 | 1-375 | 1 - 564 |
| | | 876430 | 779645 | 954090 | 781411 | 847952 | 796659 | 732436 | 723164 | 724619 | 682941 | 699695 | 772422 | 722195 | 847966 | 660552 | 720008 | 864344 | 661560 | 963157 |
| HBJCQ65 HBJCQ65 HBJCB83 HBJCB83 HBJAP95 HBJAP95 HBJAP95 HBJAB77 HBJAB17 HBJAB17 HBJAB17 HBJAB17 HBJAB17 HBJAB16 HBJAB17 HBJAB17 HBJAB16 | | 9508 | 9509 | 9510 | 9511 | 9512 | 9513 | 9514 | 9515 | 9516 | 9517 | 9518 | 9519 | 9520 | 9521 | 9522 | 9523 | 9524 | 9525 | 9526 |
| | | HBJCQ65 | HBJCJ82 | HBJCE07 | HBJCB83 | HBJBR04 | HBJAP95 | HBJA055 | HBJAJ49 | HBJAI50 | HBJAH27 | HBJAE32 | HBJAB77 | HBJAB49 | HBJAB28 | HBJAB15 | HBDAE47 | HBDAD64 | HBDAD16 | HBCC010 |

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|---|---|--|--|--|---|----------|-------------------------|-----------------------------------|---|--|---|---|--|---|---|---|---|---|---|---|---|---|---|---|---|
| H59931, H67432, Z70280, AC003100, AC006998, AC003072, AC005154, AL022154, and AC003034. | AI672457, AA810483, AI269914, AI949892, AI198001, W86511, AW237699, AI863804, and AI525426. | AI078128, AI871101, AI274339, AA035467, AA419038, AI479404, AI677732, AW190726, AI198435, AA031617, AI283200, AI983969, AA411122, AI093316, AI361637, AI769275, AI560217, AA961077, AI291805, AI291474, AA915909, AA035466, AA046943, AI857306, AA661657, AA427407, AA725194, AA423792, AW070742, AW051192, AI275083, AW293787, AW457102, AA250787, AA031475, NO3212, HOLTER AWA57102, AA250787, AA031475, NO321475, NO3214 | 111 102102, 1212,00101, AAO21470, 1722012, 117173, AW 147470, 1171000, and AAO14548. | AA280776, AL137689, AF111173, and AF100172 | AA464030, AA434281, AA410674, D78694, AI860751, AI628190, AA568168, AI097605, AI199027, D78920, AA227411, AI200649, AI199599, AI935345, AA525327, AI475176, AI910441, AI923458, AI247190, AI687257, AI923451, AI541551, AI699991, AI653151, and AW438934. | H40594. | AA677024, and AI022041. | AI566483, AA417101, and AL021368. | AA486018, AA005417, H56995, AL031666, and AF144233. | AA419501, AA969163, AA157911, AA410996, AW452982, AW339854, and H14554 | AA826669, AA287363, AI809776, AA470582, AA516190, AA505108, AA838091, AI680323, AA847361 797054 AI 040604 A CO05404 A CO0452 A CO05200 A CO0452 | XXX717501, Z77034, ALV423034, ACV03464, ACV044233, ACV03288, ACV04019, ALV49/37, ACV04884 ACV04814 ACV04014 ACV04305 ACV04805 ALV3315 AVV131318 | U95742, AL049766, AL009181, AL023553, AC016830, AC002350, AC007216, AC000652 | Z85986, AC007676, AC005562, AC016027, AC005822, AL031848, AL050318, AL031662, | AC006088, AC003950, AC005086, AC004099, AB001523, AP000694, AC004841, AC010205, | AC004765, AC005280, AC004813, AL109984, AC005409, AC003101, AC004678, AC004812, | AP000514, AL034429, AC004874, AL031602, AC006487, AJ009610, AC004596, AL136295, | AC002314, AC00/227, AL117694, AP001060, AC004125, AL031587, AL133246, AC005057, | AF001548, AF000512, AC005004, AC005088, Z83844, AL031311, U91323, AC007686, | AF001549, AL022165, AF030453, Z98941, Z84469, AC004383, AC005914, Z93016, AC005516, | AL049369, AC006039, 293017, 283847, AC005837, AC004913, AF217403, AC005015, | AC004662, AC005081, AC00/193, AC008039, AC003982, AC005058, AC006130, M89651, | AC0083/2, AF043945, AC005696, AL096701, AL024474, AC002312, AL031276, AC004531, | AI.049758. AI.035249 AI.023807 AD000501, AC002040, AI.124720, AL090712, AL031983, | AC008101, AL035086, AC006430, AC002425, Z86090, AC005082, AC007899, AC005049, |
| 15 - 582 | 15 - 593 | 15 - 604 | 15 - 529 | 15 - 581 | 15 - 1009 | 15 - 539 | 15 - 602 | 15 - 531 | 15 - 772 | 15 - 909 | 15 - 363 | - | | | | | | | | | | | | | |
| 1 - 568 | 1 - 579 | 1 - 590 | 1-515 | 1 - 567 | 1 - 995 | 1 - 525 | 1 - 588 | 1 - 517 | 1 - 758 | 1 - 895 | 1 - 349 | | | | | | | | | | | | | | |
| 930956 | 965869 | 938319 | 975256 | 958151 | 952057 | 934110 | 926761 | 969578 | 848322 | 848325 | 573919 | | | | - | | | | | | | | | | |
| 9527 | 9528 | 9529 | 9530 | 9531 | 9532 | 9533 | 9534 | 9535 | 9536 | 9537 | 9538 | | | | | | | | | | _ | | 751 | | |
| HBCCJ05 | HBCCE11 | HBCCD06 | HBCCB51 | HBCBR08 | HBCBN51 | HBCBN06 | HBCBG04 | HBCBF12 | HBCBE57 | HBCBB22 | HBCAW52 | | | | | | | | | | | | | | |
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| AL024507, AC006064, AF207550, AL135960, AJ131016, AC006312, AC006254, AC004477, AB0223049, AC005037, AC002563, AF165926, AC007199, AC004551, AL022302, AL049795, AC007191, AC005920, AC002115, AC004650, AC004694, D84394, AC005046, AL031133, AC005808, AL035659, AC009516, Z84480, AC007666, AC005529, AL049830, AC005668, AL031229, AC008040, AC006211, AL008723, AF053356, Z82244, AL022476, AL049759, AL133163, AL031283, AL109758, AP000032, AP000103, AC004851, L78810, AL080317, AP000134, AP000212, AC004817, AC002316, AL021939, AC006141, AP000689, AC006539, and AL035458. | 1 - 323 15 - 337 AI351598, and AL021528. | | 1 - 362 15 - 376 R60453, H10942, and Z42911. | 1 - 418 15 - 432 AA594734, AA376358, N69399, AI623364, AA368329, AI733856, AI246386, AA077619. | AA133013, AI246061, AA878492, AI689198, AA299422, AI268019, T53133, AI273990, | AA484022, AA211918, H73550, AW161459, AA434078, W26972, AA831638, AA614647 | AW161879, M78026, AI191343, AI811846, AI003797, AA814878, AA936548, AW268329. | AA714524, AA804334, H67064, AW069227, F33126, AA504951, T08386, AI905588, AI491765 | H24953, AI634187, AA584207, AI457313, AI280574, AA356376, AI243793, AA363027 | AA344645, AW007980, AA706202, AA226357, AW378516, AA325115, A1744933, C14966 | AA226356, AI049630, F32591, AA568399, AA058312, AI052776, AA809089, AI904840 | AA598605, H65213, AA568400, AA568416, AW378511, AW270429, AA565319, AA425924 | AI926717, AW270351, AI821420, AI697425, T47138, AA862312, AL042735, AA573693. | AI805261, AI054090, AA514450, AI469392, AI368862, AA357878, AA525331, AI718462, | AI568147, C15504, AI332615, AA363003, AA523695, AW068786, AA714512, H44944, | AA526216, AI683116, AW243793, AL033392, AC004491, AC005231, AL022326, U62293, | U63721, AC005057, AP000516, AC002375, AL031280, U02051, AC005944, AC004801, | AC007666, AL031602, AL022311, AP000696, AL022324, AP000690, AC005585, AC004662, | AJ251973, AL078638, AC007308, AL031595, AC002470, AC003030, AC005280, AC002492, | AL049646, AC006050, Z93244, AL031665, AL020993, AL049563, AL031291, AP000501, | AL109627, AC005015, AC005786, AF165926, AC006600, AF031078, AC005412, AL133243, | AL031289, AC007376, AC005317, AF030876, AC004703, AC004382, AL035455, AP000689 | AC005829, AL031007, X58139, AC009510, AC007686, U47924, AC006011, AC005011. | AL022320, AC004973, AC004865, Z94802, AC005670, AL135959, AC007707, AC007685. | AC002558, AB013139, AL031431, AB020690, AC006120, AC007358, AB000931, AC005393 | AC005908, AC000025, AC004651, AC005316, AC004760, AC003071, AC005031, AL031577. | AC000052, AC004817, AC005837, AL022721, AC007880, AC002115, U63834, Z82250, | AF043460, AC007390, AL078602, AC007263, U78027, AC004230, M84472, AL079342, | AL035682, AC005214, AL022334, AC004796, AC007444, AF051976, AC000125, AC000125, AC000125, AC000125, AC000126, AC00 |
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| | 968195 | | 783431 | 864356 | | | | | | | | | | | | | | | | | | | | | | | _ | | |
| | 9539 | 9540 | | 9542 | | | | | | | | _ | | | | | _ | | | | | | | | · | | | | |
| | HBCAT10 | HBCAS32 | HBCAQ85 | HBCAQ42 | | | | | | | | | | | | | | | | | | | | | | | | | |

| 2627 | HASCG58 HASAW90 HASAR74 | 9543 9545 9545 | 738423 789112 879377 | 1 - 489 1 - 822 1 - 1185 | 15 - 503 15 - 836 15 - 1199 | AC0065913, AP000692, AC004912, AC005747, AL035422, Z95116, AL035461, AC007543, Z84469, AP000172, Z93017, AF069291, AL035685, AC004910, AC000039, AP0000157, AC006254, AC004966, AC0049890, AC005215, AC004910, AC0003037, Z94161, AP000701, AC006211, AC004217, AL022322, Z98044, Z83846, AC005480, AC003037, Z94161, AP000701, AP0000272, AC0004217, AL022322, Z98044, Z83846, AC002483, AC003963, AL035420, AL020204, AL096791, AP000695, U3819, AC004832, Z8487, AC005216, AC003004, AF008191, AF049895, AC006130, AC00245, AC003963, AL035420, AL02726, AF035423, AC006581, AC004849, AC006522, Z85997, AD000092, AC005391, AL021653, AL049872, AD000813, AC004834, Z97630, Z81010, AC007406, AC00539, Z8244, AP000210, AF000210, AF000210, AF002397, AC006231, AL023876, AC006580, U31561, AC007160, AC005099, Z82244, AC0064752, AC000239, AL032876, AC006580, AL110658, AC006438, AC006491, AL035681, AL109865, AC00438, AC006432, AC006432, AC006411, AL0309051, AF000550, AC004988, AC006438, AC006438, AC006438, AC006438, AC006438, AC006438, AC006439, AC006439, AC006441, AL0360951, AR000550, AC004988, AC006438, AC006443, AC006444, AC006441, AL036701, AC006501, AC005082, AL049699, AC006538, and AA625541. AV30215, AA416659, AL584030, AA601356, AA140660, D78802, AN182288, and AA625541. AV30215, AA416659, AA584489, AV188742, AA581263, AA714011, AA749233, AA661313, AA687730, AA584489, AV188742, AA581364, AA190177, AA1827114, AL135357, AA584489, AV188742, AL39336, AC004615, Z96074, AC006409, AL032383, U95743, AL117329, AC005071, AL035405, AC006281, AC006329, AC006493, AC006404, AL136297, AC006497, AC006319, AC00631, AL039364, AC006244, AC006281, AC006493, AC006493, AC006493, AC006493, AC006493, AC006497, AC006319, AC00638, AC006483, AC006493, AC006493, AC006493, AC006493, AC006493, AC006493, AC006493, AC006534, AC006534, AC0065363, AC00653636, AC0065363, AC0065363, AC0065363, AC0065363, AC0065363, AC0064 |
|------|-------------------------------|----------------------|----------------------------|--------------------------------|-----------------------------------|--|
| | | | | | | AL022323, U95741, AC002984, AL121694, AC007981, AP00208, AP000557, AC005013, AC002323, U95741, AC002984, AL121694, AC007981, AP000208, AP000557, AC005013, AC002542, Z82201, AC005821, AC005899, AL117352, AL031466, AC005512, AC005261, AC004804, AP000247, Z95114, AL031767, AP000130, AC0064101, AC006459, AC003071, U96629, AL049631, Z83822, AC006162, AC005786, AC004913, AP000030, AC005562, AC005786, AC005778, AC005874, AF134471, AP000501, AC004814, AL080241, AC004837, AC007546, AL109628, Z84480, AC007052, AC006445, AL109798, AP000712, AP000134, Z98046 |

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| AC009396, AL021155, AF172277, AC004687, AC002310, AC007461, AC000378, AL079342, | AC003004, AL008/18, AL022316, AF217403, AL031311, AC005224, AC004222, AL031650, | AP000086, AL078639, AC007664, Z69710, AP000251, AB000877, AC004975, AC007792. | AL135783, AC004491, AC007406, AC005529, AC007055, AL035587, Z82203, AC002504 | AC004707, AC002350, AC005940, AL050333, Z93023, AL135744, AP000516, AC007919, | AC004859, U78027, AL035684, Z81364, Z84466, Z95115, Z85987, AC004128, AC006046, | AC004386, AC005884, AC004775, AL035422, AC005544, AC007541, AL031733, AC005088. | AL035420, Z85986, AC002091, AC009247, AC005519, AC005907, AC002558, AL139054. | AC000385, AL136295, AC007488, AP000506, AC006211, AC007225, AC000025, AC005527. | AL031431, AC005412, AC004797, AC005587, AC002369, AC004865, AC005683, AP000502, | AL050338, AF134726, AC008132, AL136504, Z75887, AC005578, AL137191, AL030995. | AF109907, AC006088, AB000882, AC002375, AL008627, AC005042, AC004659, AL021807. | AF099810, AC004125, AC005377, AC007676, AF024534, AD001527, AC002119, AC004841, | AC006333, AL109952, AC005921, AL035072, AC004694, AP000065, AC005015, AL050308, | Z98200, AC006530, AC004156, AC002486, AC004453, AL078581, AC011311, AC005919 | AC002416, AC004598, AC006241, and AF001549. | AI863136, N71484, AA318470, AA972121, AW129333, and AC007677. | |
| | | | | | | | | | | | | | | | | 15 - 484 | |
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TABLE 4

| Code | Description | Tissue | Organ | Cell Line | Disease | Vector |
|--------|-----------------------------|-------------------------------|-------------|--|---------|-------------|
| AR022 | a_Heart | a Heart | | | | |
| AR023 | a_Liver | a_Liver | | | | |
| AR024 | a_mammary gland | a_mammary gland | | | | |
| AR025 | a_Prostate | a_Prostate | | | | |
| AR026 | a_small intestine | a_small intestine | | | | |
| AR027 | a_Stomach | a_Stomach | 1 | | | |
| AR028 | Blood B cells | Blood B cells | | | *** | |
| AR029 | Blood B cells activated | Blood B cells | | | | |
| | | activated | | | | |
| AR030 | Blood B cells resting | Blood B cells | | | | |
| | | resting | <u> </u> | | | |
| AR031 | Blood T cells activated | Blood T cells | | | | |
| | | activated | | | | |
| _AR032 | Blood T cells resting | Blood T cells resting | | | | |
| AR033 | brain | brain | | | | |
| AR034 | breast | breast | | | | |
| AR035 | breast cancer | breast cancer | | | | |
| AR036 | Cell Line CAOV3 | Cell Line CAOV3 | | | | |
| _AR037 | cell line PA-1 | cell line PA-1 | | | | |
| AR038 | cell line transformed | cell line transformed | | | | |
| AR039 | colon | colon | | | | |
| AR040 | colon (9808co65R) | colon (9808co65R) | | | | |
| AR041 | colon (9809co15) | colon (9809co15) | | | | |
| AR042 | colon cancer | colon cancer | | | | |
| AR043 | colon cancer (9808co64R) | colon cancer (9808co64R) | | | | |
| AR044 | colon cancer 9809co14 | colon cancer 9809co14 | | | | |
| AR045 | corn clone 5 | corn clone 5 | | | _ | |
| AR046 | corn clone 6 | corn clone 6 | | | | |
| AR047 | corn clone2 | corn clone2 | | - | | |
| AR048 | corn clone3 | corn clone3 | | | | |
| AR049 | Corn Clone4 | Corn Clone4 | | | | |
| AR050 | Donor II B Cells 24hrs | Donor II B Cells | | · · · · · · · · · · · · · · · · · · · | | |
| | | 24hrs | | | | |
| AR051 | Donor II B Cells 72hrs | Donor II B Cells 72hrs | | | | |
| AR052 | Donor II B-Cells 24 hrs. | Donor II B-Cells 24 | | | _ | |
| | | hrs. | | | | |
| AR053 | Donor II B-Cells 72hrs | Donor II B-Cells | | | _ | |
| | | 72hrs | |] | ļ | ! |
| AR054 | Donor II Resting B Cells | Donor II Resting B | | | | |
| | _ | Cells | | | | |
| AR055 | Heart | Heart | | | | |
| AR056 | Human Lung (clonetech) | Human Lung (clonetech) | | | | |
| AR057 | Human Mammary (clontech) | Human Mammary (clontech) | | | | |
| AR058 | Human Thymus (clonetech) | Human Thymus (clonetech) | | | | |
| AR059 | Jurkat (unstimulated) | Jurkat | | | | |
| 111000 | on the (ansimulated) | (unstimulated) | | | ļ | |
| AR060 | Kidney | Kidney | | | | |
| AR061 | Liver | Liver | | 1 | | |
| AR062 | Liver (Clontech) | Liver (Clontech) | | + | | |
| AR063 | Lymphocytes chronic | Lymphocytes | | | | |
| | lymphocytic leukaemia | chronic lymphocytic leukaemia | | | | |

| AR064 | Lymphocytes diffuse large | Lymphocytes | 1 | | 1 | |
|----------------|--------------------------------------|---------------------------|---------------|--|--|--|
| 111007 | B cell lymphoma | diffuse large B cell | | | 1 | |
| | | lymphoma | | | | ļ |
| AR065 | Lymphocytes follicular | Lymphocytes | | | | |
| į. | lymphoma | follicular lymphoma | | | | |
| AR066 | normal breast | normal breast | <u> </u> | | | |
| AR067 | Normal Ovarian | Normal Ovarian | | | | |
| | (4004901) | (4004901) | | | - | |
| AR068 | Normal Ovary 9508G045 | Normal Ovary | | | | <u> </u> |
| | | 9508G045 | | 1 | | |
| AR069 | Normal Ovary 9701G208 | Normal Ovary | | | 1 | |
| | | 9701G208 | | | | |
| AR070 | Normal Ovary 9806G005 | Normal Ovary | | | | |
| | | 9806G005 | | İ | İ | 1 |
| AR071 | Ovarian Cancer | Ovarian Cancer | | | | |
| AR072 | Ovarian Cancer | Ovarian Cancer | 1 | 1 | | |
| | (9702G001) | (9702G001) | | | | |
| AR073 | Ovarian Cancer | Ovarian Cancer | | | | |
| | (9707G029) | (9707G029) | | | | 1 |
| AR074 | Ovarian Cancer | Ovarian Cancer | | | | T |
| \ | (9804G011) | (9804G011) | | <u> </u> | | |
| AR075 | Ovarian Cancer | Ovarian Cancer | | | | |
| 17076 | (9806G019) | (9806G019) | | | | |
| AR076 | Ovarian Cancer | Ovarian Cancer | | | | |
| AR077 | (9807G017) | (9807G017) | <u> </u> | ! | | ļ |
| AKUII | Ovarian Cancer (9809G001) | Ovarian Cancer | | | | |
| AR078 | ovarian cancer 15799 | (9809G001) | | ļ | - | |
| ARU/6 | Ovarian cancer 13/99 | ovarian cancer 15799 | | ľ | } | |
| AR079 | Ovarian Cancer | Ovarian Cancer | | | ļ | |
| AROTS | 17717AID | 17717AID | | J | | |
| AR080 | Ovarian Cancer | Ovarian Cancer | - | | | |
| 7111000 | 4004664B1 | 4004664B1 | | | | |
| AR081 | Ovarian Cancer | Ovarian Cancer | | | - | |
| 1111001 | 4005315A1 | 4005315A1 | | | 1 | |
| AR082 | ovarian cancer 94127303 | ovarian cancer | | | | |
| | | 94127303 | • | ĺ | | |
| AR083 | Ovarian Cancer 96069304 | Ovarian Cancer | | | <u> </u> | |
| | | 96069304 | | | | |
| AR084 | Ovarian Cancer 9707G029 | Ovarian Cancer | | | <u> </u> | |
| | | 9707G029 | | | | |
| AR085 | Ovarian Cancer 9807G045 | Ovarian Cancer | | | | |
| | | 9807G045 | | | | |
| AR086 | ovarian cancer 9809G001 | ovarian cancer | | | | |
| | | 9809G001 | | | | |
| AR087 | Ovarian Cancer | Ovarian Cancer | | | | |
| A 70,000 | 9905C032RC | 9905C032RC | | | | |
| AR088 | Ovarian cancer 9907 C00 | Ovarian cancer 9907 | | ļ | 1 |] |
| 09097 | 3rd Prostate | C00 3rd | | | ļ <u></u> | |
| AR089 AR090 | Prostate (clonetech) | Prostate (claratech) | | | | |
| AR090 AR091 | Prostate (clonetech) prostate cancer | Prostate (clonetech) | | | | |
| AR091 AR092 | prostate cancer #15176 | prostate cancer | | | | |
| ANU34 | prostate cancer #13176 | prostate cancer #15176 | | |] |] |
| AR093 | prostate cancer #15509 | prostate cancer | | | | |
| 1111075 | prestate carreer #15505 | #15509 | | | | |
| AR094 | prostate cancer #15673 | prostate cancer | | | | |
| | F- 55445 CHICOI # 15075 | #15673 | | | 1 | j J |
| AR095 | Small Intestine (Clontech) | Small Intestine | | | | |
| | (0,0,1,0,0,1) | (Clontech) | | | | |
| AR096 | Spleen | Spleen | | | <u> </u> | |
| AR097 | Thymus T cells activated | Thymus T cells | | | | |
| | | | | | <u> </u> | |

| | | activated | | | | |
|----------------|---|----------------------------|--|--|--------------|------------------------|
| AR098 | Thymus T cells resting | Thymus T cells | | | | |
| | | resting | | | 1 | |
| AR099 | Tonsil | Tonsil | | | | - |
| AR100 | Tonsil geminal center | Tonsil geminal | | | | - |
| 1.77.40.1 | centroblast | center centroblast | | | _1 | |
| AR101 | Tonsil germinal center B | Tonsil germinal | | | | |
| A D 100 | cell | center B cell | | | | |
| AR102 AR103 | Tonsil lymph node | Tonsil lymph node | | | | |
| AK103 | Tonsil memory B cell | Tonsil memory B | - | _ | | |
| AR104 | Whole Brain | cell | - | <u> </u> | | |
| AR105 | Xenograft ES-2 | Whole Brain Xenograft ES-2 | | | | |
| AR106 | Xenograft SW626 | Xenograft SW626 | | - | | |
| H0004 | Human Adult Spleen | Human Adult | Calcar | | - | |
| | Taman Franciscon | Spleen | Spleen | | 1 | Uni-ZAP |
| H0013 | Human 8 Week Whole | Human 8 Week Old | Embryo | | | XR |
| | Embryo | Embryo | Lindiyo | 1 | | Uni-ZAP XR |
| H0022 | Jurkat Cells | Jurkat T-Cell Line | | | | Lambda |
| | | | ļ | | | ZAP II |
| H0025 | Human Adult Lymph | Human Adult | Lymph Node | 1 | | Lambda |
| | Node | Lymph Node | | | | ZAPII |
| H0031 | Human Placenta | Human Placenta | Placenta | | | Uni-ZAP |
| 110050 | | | | <u> </u> | 1 | XR |
| H0050 | Human Fetal Heart | Human Fetal Heart | Heart | | | Uni-ZAP |
| H0057 | Human Estal S. I | | | <u> </u> | | XR |
| 110037 | Human Fetal Spleen | | | ł | | Uni-ZAP |
| H0058 | Human Thymus Tumor | I In The The I | | | <u> </u> | XR |
| 110030 | Truman Thymus Tumor | Human Thymus Tumor | Thymus | | disease | Lambda |
| H0059 | Human Uterine Cancer | Human Uterine | T.74 | <u> </u> | | ZAPII |
| 110055 | Trainan Oternie Cancer | Cancer | Uterus | | disease | Lambda |
| H0060 | Human Macrophage | Human Macrophage | Blood | Cell Line | ļ | ZAPII |
| H0061 | Human Macrophage | Human Macrophage | Blood | Cell Line | | pBluescript |
| H0062 | Human Thymus | Human Thymus | Thymus | Cen Line | | pBluescript Uni-ZAP |
| | | 1 11/11/45 | inymus | ļ | | XR |
| H0063 | Human Thymus | Human Thymus | Thymus | | | Uni-ZAP |
| | | | | Í | ļ | XR XR |
| H0069 | Human Activated T-Cells | Activated T-Cells | Blood | Cell Line | | Uni-ZAP |
| 7700=0 | | | | | | XR |
| H0073 | Human Leiomyeloid | Human Leiomyeloid | Muscle | | disease | Uni-ZAP |
| 110074 | Carcinoma | Carcinoma | | | | XR |
| H0074 | Human Platelets | Human Platelets | Blood | Cell Line | | Uni-ZAP |
| H0075 | Human Activated T-Cells | A 41 - 1 - 1 - 1 - 1 | | | | XR |
| 110075 | (II) | Activated T-Cells | Blood | Cell Line | | Uni-ZAP |
| H0083 | HUMAN JURKAT | Jurkat Cells | | | | XR |
| 110003 | MEMBRANE BOUND | Jurkat Cens | | | | Uni-ZAP |
| | POLYSOMES | | | | | XR |
| H0087 | Human Thymus | Human Thymus | | | | -Dli |
| H0090 | Human T-Cell Lymphoma | T-Cell Lymphoma | T-Cell | | disease | pBluescript |
| | 3 1 | | 1 - COII | | disease | Uni-ZAP XR |
| H0108 | Human Adult Lymph | Human Adult | Lymph Node | | | Uni-ZAP |
| | Node, subtracted | Lymph Node | , , | | | XR |
| H0109 | Human Macrophage, | Macrophage | Blood | Cell Line | | pBluescript |
| | subtracted | | | | | poraescript |
| H0116 | Human Thymus Tumor, | Human Thymus | Thymus | | W | pBluescript |
| | subtracted | Tumor | | , | | Parassirpi |
| H0128 | Jurkat cells, thiouridine | Jurkat Cells | | | | Uni-ZAP |
| 170156 | activated | | | | | XR |
| H0129 | Jurkat cells, thiouridine activated, fract II | Jurkat Cells | | | | Uni-ZAP |
| 1 | | | | | | |

| H0134 | Raji Cells, cyclohexamide treated | Cyclohexamide Treated Cem, Jurkat, Raji, and Supt | Blood | Cell Line | Uni-ZAP XR |
|-------|--|---|------------|-----------|-------------------------|
| H0139 | Activated T-Cells, 4 hrs. | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0140 | Activated T-Cells, 8 hrs. | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0141 | Activated T-Cells, 12 hrs. | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0158 | Activated T-Cells, 4 hrs., ligation 2 | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0159 | Activated T-Cells, 8 hrs., ligation 2 | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0160 | Activated T-Cells, 12 hrs., ligation 2 | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0161 | Activated T-Cells, 24 hrs., ligation 2 | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0167 | Activated T-Cells, 24 hrs. | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0179 | Human Neutrophil | Human Neutrophil | Blood | Cell Line | Uni-ZAP XR |
| H0185 | Activated T-Cell labeled with 4-thioluri | T-Cells | Blood | Cell Line | Lambda ZAP II |
| H0186 | Activated T-Cell | T-Cells | Blood | Cell Line | Lambda ZAP II |
| H0187 | Resting T-Cell | T-Cells | Blood | Cell Line | Lambda |
| H0189 | Human Resting Macrophage | Human Macrophage/Monoc ytes | Blood | Cell Line | ZAP II Uni-ZAP XR |
| H0191 | Human Activated Macrophage (LPS), thiour | Human Macrophage/Monoc ytes | Blood | Cell Line | Uni-ZAP XR |
| H0202 | Jurkat Cells, cyclohexamide treated, subtraction | Cyclohexamide Treated Cem, Jurkat, Raji, and Supt | Blood | Cell Line | Uni-ZAP XR |
| H0203 | Jurkat Cells, cyclohexamide treated, dif | Cyclohexamide Treated Cem, Jurkat, Raji, and Supt | Blood | Cell Line | Uni-ZAP XR |
| H0218 | Activated T-Cells, 0hrs, subtracted | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0219 | Activated T-Cells, Ohrs, differentially expressed | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0220 | Activated T-Cells, 4 hrs, subtracted | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0221 | Activated T-Cells, 4 hrs, differentially expressed | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0222 | Activated T-Cells, 8 hrs, subtracted | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0223 | Activated T-Cells, 8 hrs, differentially expressed | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0224 | Activated T-Cells, 12 hrs, subtracted | Activated T-Cells | Blood | Cell Line | Uni-ZAP XR |
| H0225 | Activated T-Cells, 12hrs, differentially expressed | Activated T-Cells | Blood | Cell Line | Uni-ZAP |
| H0250 | Human Activated Monocytes | Human Monocytes | | | XR Uni-ZAP XR |
| H0253 | Human adult testis, large inserts | Human Adult Testis | Testis | | Uni-ZAP XR |
| H0254 | Breast Lymph node cDNA library | Breast Lymph Node | Lymph Node | | Uni-ZAP XR |
| H0255 | breast lymph node CDNA | Breast Lymph Node | Lymph Node | | Lambda |

| | library | Τ | T | 1 | | ZAPII |
|-------|--|---|-------------------|-----------|---------------------------------------|---------------------|
| H0261 | H. cerebellum, Enzyme subtracted | Human Cerebellum | Brain | | | Uni-ZAP XR |
| H0264 | human tonsils | Human Tonsil | Tonsil | | | Uni-ZAP |
| H0265 | Activated T-Cell (12hs)/Thiouridine labelledEco | T-Cells | Blood | Cell Line | | XR Uni-ZAP XR |
| H0271 | Human Neutrophil, Activated | Human Neutrophil - Activated | Blood | Cell Line | | Uni-ZAP XR |
| H0272 | HUMAN TONSILS, FRACTION 2 | Human Tonsil | Tonsil | | | Uni-ZAP XR |
| H0274 | Human Adult Spleen, fractionII | Human Adult Spleen | Spleen | | | Uni-ZAP XR |
| H0300 | CD34 positive cells (Cord Blood) | CD34 Positive Cells | Cord Blood | | | ZAP Express |
| H0305 | CD34 positive cells (Cord Blood) | CD34 Positive Cells | Cord Blood | | | ZAP Express |
| H0306 | CD34 depleted Buffy Coat (Cord Blood) | CD34 Depleted Buffy Coat (Cord Blood) | Cord Blood | | | ZAP Express |
| H0318 | HUMAN B CELL LYMPHOMA | Human B Cell Lymphoma | Lymph Node | | disease | Uni-ZAP XR |
| H0341 | Bone Marrow Cell Line (RS4;11) | Bone Marrow Cell Line RS4;11 | Bone Marrow | Cell Line | | Uni-ZAP XR |
| H0354 | Human Leukocytes | Human Leukocytes | Blood | Cell Line | | pCMVSport |
| H0369 | H. Atrophic Endometrium | Atrophic Endometrium and myometrium | | | | Uni-ZAP XR |
| H0370 | H. Lymph node breast Cancer | Lymph node with Met. Breast Cancer | | | disease | Uni-ZAP XR |
| H0376 | Human Spleen | Human Adult Spleen | Spleen | | | pCMVSport |
| H0393 | Fetal Liver, subtraction II | Human Fetal Liver | Liver | | | pBluescript |
| H0402 | CD34 depleted Buffy Coat (Cord Blood), re-excision | CD34 Depleted Buffy Coat (Cord Blood) | Cord Blood | | 1 | ZAP Express |
| H0416 | Human Neutrophils, Activated, re-excision | Human Neutrophil - Activated | Blood | Cell Line | | pBluescript |
| H0421 | Human Bone Marrow, re- excision | Bone Marrow | | | ···· | pBluescript |
| H0422 | T-Cell PHA 16 hrs | T-Cells | Blood | Cell Line | | pSport1 |
| H0423 | T-Cell PHA 24 hrs | T-Cells | Blood | Cell Line | , , , , , , , , , , , , , , , , , , , | pSport1 |
| H0436 | Resting T-Cell Library,II | T-Cells | Blood | Cell Line | | pSport1 |
| H0439 | Human Eosinophils | Eosinophils | | | | pBluescript |
| H0444 | Spleen metastic melanoma | Spleen, Metastic malignant melanoma | Spleen | | disease | pSport1 |
| H0445 | Spleen, Chronic lymphocytic leukemia | Human Spleen, CLL | Spleen | | disease | pSport1 |
| H0457 | Human Eosinophils | Human Eosinophils | | | | pSport1 |
| H0477 | Human Tonsil, Lib 3 | Human Tonsil | Tonsil | | | pSport1 |
| H0478 | Salivary Gland, Lib 2 | Human Salivary Gland | Salivary gland | | | pSport1 |
| H0485 | Hodgkin"s Lymphoma I | Hodgkin"s Lymphoma I | | | disease | pCMVSport 2.0 |
| H0486 | Hodgkin"s Lymphoma II | Hodgkin"s Lymphoma II | | | disease | pCMVSport 2.0 |
| H0487 | Human Tonsils, lib I | Human Tonsils | | | | pCMVSport 2.0 |
| H0488 | Human Tonsils, Lib 2 | Human Tonsils | | | _ | pCMVSport |

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|----------|---|--|-------------|--------------|---------|------------------|
| H0494 | Keratinocyte | Keratinocyte | | | | pCMVSport |
| H0506 | Ulcerative Colitis | Colon | Colon | | | pSport1 |
| H0510 | Human Liver, normal | Human Liver, normal, Patient #8 | Liver | | | pCMVSport |
| H0518 | pBMC stimulated w/ poly I/C | pBMC stimulated | | | | 2.0 pCMVSport |
| H0521 | Primary Dendritic Cells, | with poly I/C Primary Dendritic cells | | | | pCMVSport |
| H0522 | Primary Dendritic cells, frac 2 | Primary Dendritic cells | | | | 2.0 pCMVSport |
| H0524 | Primary Dendritic Cells, CapFinder, frac 2 | Primary Dendritic cells | | | | 3.0 pSport1 |
| H0529 | Myoloid Progenitor Cell Line | TF-1 Cell Line; Myoloid progenitor cell line | | | | pCMVSport 3.0 |
| H0537 | H. Primary Dendritic Cells,lib 3 | Primary Dendritic cells | | | | pCMVSport |
| H0542 | T Cell helper I | Helper T cell | 4.5.6. | | | pCMVSport |
| H0543 | T cell helper II | Helper T cell | | | | pCMVSport |
| H0545 | Human endometrial stromal cells-treated with progesterone | Human endometrial stromal cells-treated with proge | | | | pCMVSport 3.0 |
| H0551 | Human Thymus Stromal Cells | Human Thymus Stromal Cells | | | | pCMVSport |
| H0552 | Signal trap,Femur Bone Marrow,pooled | Femur Bone marrow, pooled from 8 male/female | | | | Other |
| H0556 | Activated T- cell(12h)/Thiouridine-re- excision | T-Cells | Blood | Cell Line | | Uni-ZAP XR |
| H0560 | KMH2 | КМН2 | | | | pCMVSport 3.0 |
| H0575 | Human Adult Pulmonary;re-excision | Human Adult Pulmonary | Lung | | | Uni-ZAP XR |
| H0576 | Resting T-Cell; re- excision | · T-Cells | Blood | Cell Line | | Lambda ZAP II |
| H0578 | Human Fetal Thymus | Fetal Thymus | Thymus | | | pSport1 |
| H0580 | Dendritic cells, pooled | Pooled dendritic cells | | | | pCMVSport 3.0 |
| H0581 | Human Bone Marrow, treated | Human Bone Marrow | Bone Marrow | | | pCMVSport |
| H0583 | B Cell lymphoma | B Cell Lymphoma | B Cell | | disease | pCMVSport 3.0 |
| H0584 | Activated T-cells, 24 hrs,re-excision | Activated T-Cells | Blood | Cell Line | | Uni-ZAP XR |
| H0585 | Activated T-Cells,12 hrs,re-excision | Activated T-Cells | Blood | Cell Line | | Uni-ZAP XR |
| H0589 | CD34 positive cells (cord blood),re-ex | CD34 Positive Cells | Cord Blood | | | ZAP Express |
| H0591 | Human T-cell lymphoma;re-excision | T-Cell Lymphoma | T-Cell | | disease | Uni-ZAP XR |
| H0606 | Human Primary Breast Cancer;re-excision | Human Primary Breast Cancer | Breast | | disease | Uni-ZAP XR |
| H0607 | H.Leukocytes, normalized cot 50A3 | H.Leukocytes | | | | pCMVSport |
| H0608 | H. Leukocytes, control | H.Leukocytes | | | | pCMVSport |
| H0609 | H. Leukocytes, | H.Leukocytes | | | | pCMVSport |

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|-------|---|--|---------------------|-----------|---------------------------------------|------------------|
| | normalized cot > 500A | | | | | 1 |
| H0610 | H. Leukocytes, normalized cot 5A | H.Leukocytes | | | | pCMVSport |
| H0611 | | III I and a second | - | + | | |
| HUGII | H. Leukocytes, normalized cot 500 B | H.Leukocytes | | | | pCMVSport |
| H0612 | H.Leukocytes, normalized cot 50 B | H.Leukocytes | | | | pCMVSport |
| H0614 | H. Leukocytes, normalized cot 500 A | H.Leukocytes | | | | pCMVSport |
| H0625 | Ku 812F Basophils Line | Ku 812F Basophils | | | | pSport1 |
| H0635 | Human Activated T-Cells, re-excision. | Activated T-Cells | Blood | Cell Line | | Uni-ZAP XR |
| H0637 | Dendritic Cells From CD34 Cells | Dentritic cells from CD34 cells | | | | pSport1 |
| H0638 | CD40 activated monocyte dendridic cells | CD40 activated monocyte dendridic cells | | | | pSport1 |
| H0641 | LPS activated derived dendritic cells | LPS activated monocyte derived dendritic cells | | | | pSport1 |
| H0650 | B-Cells | B-Cells | | | | pCMVSport 3.0 |
| H0653 | Stromal Cells | Stromal Cells | | | | pSport1 |
| H0656 | B-cells (unstimulated) | B-cells (unstimulated) | | | | pSport1 |
| H0657 | B-cells (stimulated) | B-cells (stimulated) | | | | pSport1 |
| H0659 | Ovary, Cancer (15395A1F): Grade II Papillary Carcinoma | Grade II Papillary Carcinoma, Ovary | Ovary | | disease | pSport1 |
| H0660 | Ovary, Cancer: (15799A1F) Poorly differentiated carcinoma | Poorly differentiated carcinoma, ovary | | | disease | pSport1 |
| H0673 | Human Prostate Cancer, Stage B2; re-excision | Human Prostate Cancer, stage B2 | Prostate | | | Uni-ZAP XR |
| H0677 | TNFR degenerate oligo | B-Cells | | | | PCRII |
| H0679 | screened clones from Tonsil library | Human Tonsils | | | · | Other |
| S0002 | Monocyte activated | Monocyte-activated | blood | Cell Line | | Uni-ZAP XR |
| S0026 | Stromal cell TF274 | stromal cell | Bone marrow | Cell Line | | Uni-ZAP XR |
| S0027 | Smooth muscle, serum treated | Smooth muscle | Pulmanary artery | Cell Line | | Uni-ZAP XR |
| S0031 | Spinal cord | Spinal cord | spinal cord | | | Uni-ZAP XR |
| S0052 | neutrophils control | human neutrophils | blood | Cell Line | | Uni-ZAP XR |
| S0053 | Neutrophils IL-1 and LPS induced | human neutrophil induced | blood | Cell Line | | Uni-ZAP XR |
| S0114 | Anergic T-cell | Anergic T-cell | | Cell Line | W. | Uni-ZAP XR |
| S0116 | Bone marrow | Bone marrow | Bone marrow | | V-1W | Uni-ZAP XR |
| S0134 | Apoptotic T-cell | apoptotic cells | | Cell Line | · | Uni-ZAP XR |
| S0140 | eosinophil-IL5 induced | eosinophil | lung | Cell Line | · · · · · · · · · · · · · · · · · · · | Uni-ZAP XR |
| S0142 | Macrophage-oxLDL | macrophage- oxidized LDL treated | blood | Cell Line | | Uni-ZAP XR |
| S0144 | Macrophage (GM-CSF treated) | Macrophage (GM- CSF treated) | | | | Uni-ZAP XR |

| S0180 | Bone Marrow Stroma, | Bone Marrow | | | disease | Uni-ZAP |
|-------|--|--|--------------------|-----------|---------|--------------------|
| | TNF&LPS ind | Stroma, TNF & LPS induced | | | | XR |
| S0182 | Human B Cell 8866 | Human B- Cell 8866 | | | | Uni-ZAP XR |
| S0196 | Synovial IL-1/TNF stimulated | Synovial Fibroblasts | | | | pSport1 |
| S0212 | Bone Marrow Stromal Cell, untreated | Bone Marrow Stromal Cell,untreated | | | | pSport1 |
| S0216 | Neutrophils IL-1 and LPS induced | human neutrophil induced | blood | Cell Line | | Uni-ZAP XR |
| S0218 | Apoptotic T-cell, re- excision | apoptotic cells | | Cell Line | | Uni-ZAP XR |
| S0222 | H. Frontal cortex,epileptic;re- excision | H. Brain, Frontal Cortex, Epileptic | Brain | | disease | Uni-ZAP XR |
| S0278 | H Macrophage (GM-CSF treated), re-excision | Macrophage (GM- CSF treated) | | | | Uni-ZAP XR |
| S0282 | Brain Frontal Cortex, re- excision | Brain frontal cortex | Brain | | | Lambda ZAP II |
| S0292 | Osteoarthritis (OA-4) | Human Osteoarthritic Cartilage | Bone | | disease | pSport1 |
| S0298 | Bone marrow stroma,treated | Bone marrow stroma,treatedSB | Bone marrow | | | pSport1 |
| S0308 | Spleen/normal | Spleen normal | | | | pSport1 |
| S0314 | Human osteoarthritis;fraction I | Human osteoarthritic cartilage | | | disease | pSport1 |
| S0344 | Macrophage-oxLDL; re- excision | macrophage- oxidized LDL treated | blood | Cell Line | | Uni-ZAP XR |
| S0358 | Colon Normal III | Colon Normal | Colon | | | pSport1 |
| S0418 | CHME Cell Line;treated 5 hrs | CHME Cell Line; treated | | | | pCMVSport |
| S0426 | Monocyte activated; re- excision | Monocyte-activated | blood | Cell Line | | Uni-ZAP XR |
| S0428 | Neutrophils control; re- excision | human neutrophils | blood | Cell Line | | Uni-ZAP XR |
| S0442 | Colon Normal | Colon Normal | | | | pSport1 |
| S0452 | Thymus | Thymus | | | | pSport1 |
| S0474 | Human blood platelets | Platelets | Blood platelets | | | Other |
| S3016 | Basophil | Basophil | | | | Uni-ZAP XR |
| S3018 | TH1 cells | TH1 cells | 4 | | | Uni-ZAP XR |
| S3020 | TH2 cells | TH2 cells | | | ··· | Uni-ZAP XR |
| S6028 | Human Manic Depression Tissue | Human Manic depression tissue | Brain | | disease | Uni-ZAP XR |
| T0002 | Activated T-cells | Activated T-Cell, PBL fraction | Blood | Cell Line | | pBluescript SK- |
| T0041 | Jurkat T-cell G1 phase | Jurkat T-cell | | | | pBluescript SK- |
| T0042 | Jurkat T-Cell, S phase | Jurkat T-Cell Line | | | | pBluescript SK- |
| T0071 | Human Bone Marrow | Human Bone Marrow | | | | pBluescript SK- |
| L0002 | Atrium cDNA library Human heart | | | | | |

| L0004 Clonfech Huran aceta | L0004 | ClarTeck III 1005 | | | | | |
|--|----------|---------------------------|--|--------------|--|-------------|--------------|
| DeliyA+ mRNA (#6572) | | | | | | | |
| L0021 Human dutivated eduntitic cell mRNA | L0005 | | | | | | |
| L0020 | | | | | | | |
| Cell mRNA | | | | | | | |
| L0021 | L0020 | Human activated dendritic | | | | | |
| L0036 | | cell mRNA | | i | | | |
| L0036 | L0021 | Human adult (K.Okubo) | | | | | <u> </u> |
| 1,0040 | L0036 | | | | | | |
| L0040 Human colon mucosa L0041 Human city Human city Human city L0055 Human promyelocyte L0051 Human promyelocyte L0055 Human promyelocyte L0056 Human thyman SYTH II L0057 Human thyman SYTH II L0058 Liver HepG2 cell line. L0070 Selected chromosome 21 CDNA library L0109 Human brain cDNA Drain L0104 Human placenta cDNA Drain L0143 Human placenta dDNA Drain L0157 Human fetal brain L01 | | | | | | | |
| L0051 | 1.0040 | | | | | | ļ |
| L0051 Human mRNA (Tripodis and Ragoussis) | | | | ļ | | ļ | |
| L0051 | L0041 | | | | | | |
| And Ragoussis | 1.0051 | | | | | | |
| L0055 | L0051 | | - | ı | | | |
| L0060 | | | | į . | | | i |
| L0070 L0070 Selected chromosome 21 | | Human promyelocyte | | | | | |
| L0070 L0070 Selected chromosome 21 | L0060 | Human thymus NSTH II | | | | | |
| L0070 Selected chromosome 21 cDNA library L0109 Human brain cDNA brain L0142 Human placenta cDNA placenta CTrujiwara) L0143 Human placenta polyA+ (TFujiwara) L0157 Human feat brain (Fujiwara) L0163 Human feat brain (Fujiwara) L0171 Human teat brain (Fujiwara) L0171 Human mewborn melanocytes (T.Vogt) L0177 Human newborn melanocytes (T.Vogt) L0239 Homo sapiens brain fetus brain C3- A11N L0307 Human E8CASS breast adenocarcinoma E8CASS; variant of OCI LY8-C3P COCI LY8-C3P L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of MCF7 L0362 Stratagene ovary (#937217) SK L0363 NCLCGAP_GC2 germ cell tumor Bluescript SK L0365 NCLCGAP_C2C germ cell tumor Bluescript SK Stratagene schizo brain Schizophrenic brain Shitagene schizo brain Schizophrenic brain Shitagene schizo brain Schizophrenic brain Shitagene schizo brain Schizophrenic brain SK SK SK SK SK SK SK S | L0065 | | | | | | |
| CDNA library | | Selected chromosome 21 | | | | ļ | |
| L0199 Human brain cDNA Placenta Plac | 1 20070 | | | | | | |
| L0142 | T 0100 | | | | | | |
| L0143 | | | | | | | |
| L0143 | L0142 | | placenta | 1 | | | |
| CTFujiwara CTF | <u> </u> | | | <u> </u> | L | | |
| L0157 | L0143 | | placenta | | | | |
| L0163 Human heart cDNA (YNakamura) L0171 Human lung adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 adenocarcinoma A549 Clonetics Corp. (San Diego, CA) Strain #68 and 2486 | | | | | _1 | | |
| L0163 | L0157 | Human fetal brain | | brain | | | |
| LO171 Human lung adenocarcinoma A549 adenocarcinoma A549 | | (TFujiwara) | | | 1 | | |
| LO171 Human lung adenocarcinoma A549 adenocarcinoma A549 | L0163 | Human heart cDNA | | heart | | | |
| L0171 | } | | | l nour | | İ | |
| L0177 Human newborn Clonetics Corp. (San Diego, CA) Strain #68 Stratagene ovary (#937217) Clonetics Corp. (San Diego, CA) Clonetics L0171 | | lung | | A 540 | | |
| L0177 | | | | | A349 | l | |
| Melanocytes (T.Vogt) Corp. (San Diego, CA) Strain #68 and 2486 | E0177 | | adenocaremonia | | | ļ | |
| L0239 Homo sapiens brain fetus Brain | LO1// | 1 | 1 | l | | | |
| L0239 Homo sapiens brain fetus Brain | | metanocytes (1.vogt) | | | | | |
| L0239 Homo sapiens brain fetus Brain CA | 1 | | | | | | |
| L0239 Homo sapiens brain fetus | İ | 1 | | | | | |
| L0239 Homo sapiens brain fetus Brain C3- C3- A11N; clonally related variant of OCI LY8-C3P L0309 Human E8CASS Breast adenocarcinoma E8CASS; variant of MCF7 BA, M13- derived C1361 Stratagene ovarian cancer (#937217) SK- C1362 Stratagene ovarian cancer (#937219) C1363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- C1366 Stratagene schizo brain S-11 frontal lobe SK- Bluescript Bluescript Bluescript Bluescript Bluescript Bluescript Bluescript Bluescript Blu |] | | i | | CA) | | |
| L039 Homo sapiens brain fetus Brain C3- A11N; Clonally related Variant of OCI LY8-C3P L0309 Human E8CASS Breast Adenocarcinoma BA, M13- Gerived Bluescript SK- L0361 Stratagene ovarian cancer (#937217) Stratagene ovarian cancer (#937219) Stratagene ovarian cancer (#937219) Bluescript SK- L0365 NCI_CGAP_Phe1 Pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain S-11 frontal lobe SK- Bluescript SK- Bluescript SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript SK- Bluescript | 1 | 1 | | | strain #68 | | |
| L0307 Human C3-A11N C3-A11N; clonally related variant of OCI LY8-C3P | | | | | and 2486 | | |
| A11N; clonally related variant of OCI LY8-C3P | L0239 | Homo sapiens brain fetus | brain | | | | - W. |
| L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of MCF7 | L0307 | Human C3-A11N | | | C3- | | |
| L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of OCI LY8-C3P L0352 Normalized infant brain, Bento Soares L0361 Stratagene ovary (#937217) L0362 Stratagene ovarian cancer (#937219) L0363 NCI_CGAP_GC2 germ cell tumor L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain S11 Schwannoma tumor L0367 NCI_CGAP_Sch1 Schwannoma tumor L0368 L0369 L0369 Schwannoma tumor L0369 L0369 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor L0360 Schwannoma tumor | | | | | | | |
| L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of MCF7 L0352 Normalized infant brain, Bento Soares L0361 Stratagene ovary (#937217) L0362 Stratagene ovarian cancer (#937219) L0363 NCI_CGAP_GC2 germ cell tumor L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain S11 Schwannoma tumor L0367 NCI_CGAP_Sch1 Schwannoma tumor RESCASS; variant of MCF7 BA, M13-derived BA, M13-derived Bluescript SK. Bluescript SK. Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript | | | ł | | | | |
| L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of MCF7 | | <u>.</u> | | I | | | |
| L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of MCF7 | | | | | | | |
| L0309 Human E8CASS breast adenocarcinoma E8CASS; variant of MCF7 | | | | | | | |
| L0309 Human E8CASS breast adenocarcinoma L0352 Normalized infant brain, Bento Soares L0361 Stratagene ovary (#937217) L0362 Stratagene ovarian cancer (#937219) L0363 NCI_CGAP_GC2 germ cell tumor L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain S11 Schwannoma tumor Schwannoma tumor E8CASS; variant of MCF7 BA, M13-derived Bluescript SK Bluescript SK Bluescript SK- | | | | | | | |
| L0352 Normalized infant brain, Bento Soares | 1 0300 | Human ESCACE | h | | | | |
| L0352 Normalized infant brain, Bento Soares Bluescript (#937217) SK L0362 Stratagene ovarian cancer (#937219) SK- L0363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain S11 Schwannoma tumor MCF7 BA, M13- derived Bluescript SK Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript | 10303 | Tullian Locass | | | | | |
| L0352 Normalized infant brain, Bento Soares L0361 Stratagene ovary (#937217) Ovary L0362 Stratagene ovarian cancer (#937219) SK- L0363 NCI_CGAP_GC2 germ cell tumor L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain S11 Schizophrenic brain S11 Schwannoma tumor BA, M13- derived Bluescript SK Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript | | | adenocarcinoma | ĺ | | | |
| Bento Soares L0361 Stratagene ovary (#937217) Stratagene ovary L0362 Stratagene ovarian cancer (#937219) Bluescript SK- Bluescript SK- L0363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- L0365 NCI_CGAP_Phe1 pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain S11 S-11 frontal lobe L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript SK- Bluescript SK- Bluescript SK- Bluescript SK- Bluescript | T 02.52 | N. I. I. C. | | | MCF7 | | |
| L0361 Stratagene ovary (#937217) L0362 Stratagene ovarian cancer (#937219) L0363 NCI_CGAP_GC2 germ cell tumor L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain S11 S-11 frontal lobe L0367 NCI_CGAP_Sch1 Schwannoma tumor Descript SK- Schwannoma tumor Descript SK- Bluescript SK- | LU332 | | | ì | | ļ | |
| (#937217) SK L0362 Stratagene ovarian cancer (#937219) Bluescript SK- L0363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- L0365 NCI_CGAP_Phe1 pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain S11 schizophrenic brain S-11 frontal lobe Bluescript SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript | 7.00.11 | | | | | | derived |
| L0362 Stratagene ovarian cancer (#937219) Bluescript SK- L0363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- L0365 NCI_CGAP_Phe1 pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain S11 schizophrenic brain S-11 frontal lobe Bluescript SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript | L0361 | | | ovary | | | Bluescript |
| L0362 Stratagene ovarian cancer (#937219) L0363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- L0365 NCI_CGAP_Phe1 pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain schizophrenic brain S11 S-11 frontal lobe SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript Bluescript SK- | | | | | | 1 | |
| (#937219) SK- L0363 NCI_CGAP_GC2 germ cell tumor Bluescript SK- L0365 NCI_CGAP_Phe1 pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain S11 schizophrenic brain S-11 frontal lobe Bluescript SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript | L0362 | Stratagene ovarian cancer | | | | | |
| L0363 NCI_CGAP_GC2 germ cell tumor L0365 NCI_CGAP_Phe1 pheochromocytoma L0366 Stratagene schizo brain Schizophrenic brain S11 Schwannoma tumor L0367 NCI_CGAP_Sch1 Schwannoma tumor | | | | | | Į. | |
| L0365 NCI_CGAP_Phe1 pheochromocytoma SK- L0366 Stratagene schizo brain S-11 frontal lobe SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript Stratagene schizo brain S-11 frontal lobe SK- Bluescript SK- Bluescript SK- Bluescript | L0363 | NCI_CGAP_GC2 | germ cell tumor | | | | |
| L0365 NCI_CGAP_Phe1 pheochromocytoma Bluescript SK- L0366 Stratagene schizo brain S-11 frontal lobe SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript Bluescript SK- Bluescript SK- Bluescript | | | J | | | i | |
| L0366 Stratagene schizo brain Schizophrenic brain Sluescript SK- | L0365 | NCL CGAP Phe1 | pheochromocytome | | | | |
| L0366 Stratagene schizo brain schizophrenic brain S11 S-11 frontal lobe SK-L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript | 20000 | | photomorpholia | | | ŀ | |
| S11 S-11 frontal lobe SK- L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript | I 0366 | Stratagene schize hasin | sobizonhei- ti | | | | |
| L0367 NCI_CGAP_Sch1 Schwannoma tumor Bluescript | T0200 | | | | | | |
| Diacompt | 1.0267 | | | | | | |
| SK- | 下030/ | INCI_COAP_SCNI | эспwannoma tumor | | 1 | | |
| | | | | | | | SK- |

| L0368 | NCI_CGAP_SS1 | synovial sarcoma | | | | Bluescript |
|---------|--|--|---------------|--------|----------|------------------------|
| L0369 | NCI_CGAP_AA1 | adrenal adenoma | adrenal gland | | | SK- Bluescript |
| | | | | | | SK- |
| L0370 | Johnston frontal cortex | pooled frontal lobe | brain | | | Bluescript SK- |
| L0375 | NCI_CGAP_Kid6 | kidney tumor | kidney | | <u> </u> | Bluescript |
| L0376 | NCI_CGAP_Lar1 | larynx | larynx | - | | SK- Bluescript |
| <u></u> | | | | | | SK- |
| L0377 | NCI_CGAP_HN2 | squamous cell carcinoma from vocal cord | larynx | | | Bluescript SK- |
| L0378 | NCI_CGAP_Lu1 | lung tumor | lung | | | Bluescript SK- |
| L0381 | NCI_CGAP_HN4 | squamous cell carcinoma | pharynx | | | Bluescript SK- |
| L0383 | NCI_CGAP_Pr24 | invasive tumor (cell line) | prostate | | | Bluescript SK- |
| L0384 | NCI_CGAP_Pr23 | prostate tumor | prostate | | | Bluescript |
| L0385 | NCI_CGAP_Gas1 | gastric tumor | stomach | | | SK- Bluescript |
| L0386 | NCI_CGAP_HN3 | squamous cell | 4 | | _ | SK- |
| | | carcinoma from base of tongue | tongue | | | Bluescript SK- |
| L0387 | NCI_CGAP_GCB0 | germinal center B- cells | tonsil | | | Bluescript SK- |
| L0389 | NCI_CGAP_HN5 | normal gingiva (cell line from primary keratinocyt | | | | Bluescript SK- |
| L0394 | H, Human adult Brain Cortex tissue | | * 1 | | | gt11 |
| L0415 | b4HB3MA Cot8-HAP-Ft | | | | | Lafmid BA |
| L0435 | Infant brain, LLNL array of Dr. M. Soares 1NIB | | | | | lafmid BA |
| L0438 | normalized infant brain cDNA | total brain | brain | | | lafmid BA |
| L0439 | Soares infant brain 1NIB | | whole brain | | | Lafmid BA |
| L0451 | N3HFLSK20 | | | | | Lafmid K |
| L0452 | Chromosome 21, K. Gardiner | | | | | Lambda |
| L0455 | Human retina cDNA randomly primed sublibrary | retina | eye | ***** | | lambda gt10 |
| L0456 | Human retina cDNA Tsp509I-cleaved sublibrary | retina | eye | | | lambda gt10 |
| L0457 | multi-tissue normalized short-fragment | multi-tissue | pooled | | | lambda gt10 |
| L0459 | Adult heart, Clontech | | | | | Lambda gt11 |
| L0462 | WATM1 | | | | | lambda gt11 |
| L0465 | TEST1, Human adult Testis tissue | | | | | lambda nm1149 |
| L0468 | HE6W . | | | | | lambda zap |
| L0471 | Human fetal heart, Lambda ZAP Express | | | | | Lambda |
| L0475 | KG1-a Lambda Zap | | ———— <u>—</u> | KG1-a | | ZAP Express Lambda Zap |
| | Express cDNA library | | | 1501-a | | Express |
| L0476 | Fetal brain, Stratagene | | | | | (Stratagene) |
| / | orani, orangono | | | | | Lambda ZAP II |

| L0480 | Stratagene cat#937212 | | | | |
|----------------|---------------------------------------|-----------------------------------|---------------------------------------|--------------|---------------------|
| 1 20-00 | (1992) | | | | Lambda |
| ĺ | (3332) | | | | ZAP, pBluescript |
| | 1 | | 1 | | SK(-) |
| L0481 | CD34+DIRECTIONAL | | | | Lambda |
| | | | | | ZAPII |
| L0483 | Human pancreatic islet | | | | Lambda |
| 7.040.5 | | | | | ZAPII |
| L0485 | STRATAGENE Human | skeletal muscle | leg muscle | | Lambda |
| | skeletal muscle cDNA | | | | ZAPII |
| L0492 | library, cat. #936215. Human Genomic | | ļ | | |
| L0493 | NCI_CGAP_Ov26 | | | | pAMP |
| 10473 | NCI_COAF_OV20 | papillary serous carcinoma | ovary | | pAMP1 |
| L0498 | NCI_CGAP_HSC3 | CD34+, T negative, | bone marrow | | |
| | 11000 | patient with chronic | Done marrow | | pAMP1 |
| | | myelogenou | | | |
| L0506 | NCI_CGAP_Br16 | lobullar carcinoma | breast | | pAMP1 |
| | | in situ | | | PANI |
| L0508 | NCI_CGAP_Lu25 | bronchioalveolar | lung | | pAMP1 |
| | | carcinoma | | | P- 3 |
| L0509 | NCI_CGAP_Lu26 | invasive | lung | | pAMP1 |
| 7.0510 | 100 | adenocarcinoma | | | |
| L0512 | NCI_CGAP_Ov36 | borderline ovarian | ovary | | pAMP1 |
| L0513 | NCI_CGAP_Ov37 | carcinoma | | | |
| | NCI_CGAP_OV37 | early stage papillary | ovary | j | pAMP1 |
| L0515 | NCI_CGAP_Ov32 | serous carcinoma papillary serous | | | |
| 20313 | Nei_edAi_ov32 | carcinoma | ovary | | pAMP1 |
| L0517 | NCI_CGAP_Pr1 | Carcinoma | ļ | | 43 (7) (6) |
| L0518 | NCI_CGAP_Pr2 | | | | pAMP10 |
| L0519 | NCI_CGAP_Pr3 | | | | pAMP10 |
| L0520 | NCI_CGAP_Alv1 | alveolar | | | pAMP10 pAMP10 |
| | | rhabdomyosarcoma | | | PANTEIO |
| L0521 | NCI_CGAP_Ew1 | Ewing"s sarcoma | | | pAMP10 |
| L0523 | NCI_CGAP_Lip2 | liposarcoma | | | pAMP10 |
| L0526 | NCI_CGAP_Pr12 | metastatic prostate | | | pAMP10 |
| | | bone lesion | | | 1 |
| L0527 | NCI_CGAP_Ov2 | ovary | | | pAMP10 |
| L0528 | NCI_CGAP_Pr5 | prostate | | | pAMP10 |
| L0529 | NCI_CGAP_Pr6 | prostate | | | pAMP10 |
| L0530 L0532 | NCI_CGAP_Pr8 | prostate | · · · · · · · · · · · · · · · · · · · | | pAMP10 |
| L0532 | NCI_CGAP_Thy1 Chromosome 7 Fetal | thyroid | | | pAMP10 |
| L0554 | Brain cDNA Library | brain | brain | , | pAMP10 |
| L0535 | NCI_CGAP_Br5 | infiltrating ductal | breast | | 43/7210 |
| | 0 0 0. 11 _ 010 | carcinoma | oreast | | pAMP10 |
| L0539 | Chromosome 7 Placental | Garcinoma | placenta | | pAMP10 |
| | cDNA Library | | pracenta | | pAMP10 |
| L0541 | NCI_CGAP_Pr7 | low-grade prostatic | prostate | | pAMP10 |
| | | neoplasia | 1 | | priivii 10 |
| L0542 | NCI_CGAP_Pr11 | normal prostatic | prostate | | pAMP10 |
| | | epithelial cells | | | 1 |
| L0543 | NCI_CGAP_Pr9 | normal prostatic | prostate | | pAMP10 |
| 10545 | NO. CO. | epithelial cells | | | |
| L0545 | NCI_CGAP_Pr4.1 | prostatic | prostate | | pAMP10 |
| | | intraepithelial | | | |
| | | neoplasia - high | | | |
| L0546 | NCI_CGAP_Pr18 | grade | mostata | | 1117 |
| L0547 | NCI_CGAP_Pr16 | stroma tumor | prostate | | pAMP10 |
| L0551 | NCI_CGAP_HN7 | normal squamous | prostate | | pAMP10 |
| | | | | | pAMP10 |

| | | epithelium, floor of | | | | T |
|-------|---|----------------------------|--------------------|----------------------------|--------------|--------------------------------|
| L0556 | NCI_CGAP_Lu34.1 | mouth large cell carcinoma | | | | |
| L0562 | Chromosome 7 HeLa cDNA Library | large cell carcinoma | lung | HeLa cell line; ATCC | | pAMP10 pAMP10 |
| L0565 | Normal Human Trabecular Bone Cells | Bone | Hip | AICC | | pBluescript |
| L0581 | Stratagene liver (#937224) | | liver | | | pBluescript |
| L0586 | HTCDL1 | | | _ | | pBluescript |
| L0588 | Stratagene endothelial cell 937223 | 1/4/1 | | | | SK(-) pBluescript |
| L0589 | Stratagene fetal retina 937202 | | | | | pBluescript |
| L0590 | Stratagene fibroblast (#937212) | | | | | SK- pBluescript |
| L0591 | Stratagene HeLa cell s3 937216 | | | | | SK- pBluescript |
| L0592 | Stratagene hNT neuron (#937233) | | | | | pBluescript |
| L0593 | Stratagene neuroepithelium (#937231) | | | | | SK- pBluescript SK- |
| L0594 | Stratagene neuroepithelium NT2RAMI 937234 | | | | | pBluescript SK- |
| L0595 | Stratagene NT2 neuronal precursor 937230 | neuroepithelial cells | brain | | | pBluescript SK- |
| L0596 | Stratagene colon (#937204) | | colon | | | pBluescript SK- |
| L0597 | Stratagene corneal stroma (#937222) | | cornea | | | pBluescript SK- |
| L0598 | Morton Fetal Cochlea | cochlea | ear | | | pBluescript SK- |
| L0599 | Stratagene lung (#937210) | | lung | | | pBluescript SK- |
| L0600 | Weizmann Olfactory Epithelium | olfactory epithelium | nose | | | pBluescript SK- |
| L0601 | Stratagene pancreas (#937208) | | pancreas | | | pBluescript SK- |
| L0602 | Pancreatic Islet | pancreatic islet | pancreas | | | pBluescript SK- |
| L0603 | Stratagene placenta (#937225) | | placenta | | | pBluescript SK- |
| L0604 | Stratagene muscle 937209 | muscle | skeletal muscle | | | pBluescript SK- |
| L0605 | Stratagene fetal spleen (#937205) | fetal spleen | spleen | | | pBluescript SK- |
| L0606 | NCI_CGAP_Lym5 | follicular lymphoma | lymph node | | | pBluescript SK- |
| L0607 | NCI_CGAP_Lym6 | mantle cell lymphoma | lymph node | | | pBluescript SK- |
| L0608 | Stratagene lung carcinoma 937218 | lung carcinoma | lung | NCI-H69 | | pBluescript SK- |
| L0612 | Schiller oligodendroglioma | oligodendroglioma | brain | | | pBluescript SK- |
| L0617 | Chromosome 22 exon | | | | | (Stratagene) pBluescriptII KS+ |
| L0619 | Chromosome 9 exon II | | | | | pBluescriptII |

| | T | | 1 | T | 110 |
|--------|--------------------|--|-----------------|--------------|-------------------------|
| L0622 | HM1 | | | | KS+ |
| L0022 | 111/11 | | | | pcDNAII (Invitrogen) |
| L0623 | HM3 | pectoral muscle | | | pcDNAII |
| | | (after mastectomy) | | | (Invitrogen) |
| L0626 | NCI_CGAP_GC1 | bulk germ cell | | | pCMV- |
| ļ | | seminoma | | | SPORT2 |
| L0629 | NCI_CGAP_Mel3 | metastatic | bowel (skin | | pCMV- |
| L0631 | NCI_CGAP_Br7 | melanoma to bowel | primary) | | SPORT4 |
| L0031 | NCI_CGAP_BI7 | | breast | | pCMV- |
| L0633 | NCI_CGAP_Lu6 | small cell carcinoma | lung | | SPORT4 pCMV- |
| | 1 | Sman con carcinoma | rung | | SPORT4 |
| L0634 | NCI_CGAP_Ov8 | serous | ovary | | pCMV- |
| | | adenocarcinoma | | | SPORT4 |
| L0635 | NCI_CGAP_PNS1 | dorsal root ganglion | peripheral | | pCMV- |
| | | | nervous | | SPORT4 |
| 1000 | Nov. GG L D. D. 50 | | system | | |
| L0637 | NCI_CGAP_Brn53 | three pooled | brain | | pCMV- |
| L0638 | NCI_CGAP_Bm35 | meningiomas tumor, 5 pooled (see | 1 | | SPORT6 |
| L0036 | NCI_COAF_BIII33 | description) | brain | | pCMV- SPORT6 |
| L0639 | NCI_CGAP_Brn52 | tumor, 5 pooled (see | brain | | pCMV- |
| 2000 | 1101_00111 _511152 | description) | Oram | | SPORT6 |
| L0640 | NCI_CGAP_Br18 | four pooled high- | breast | | pCMV- |
| | | grade tumors, | | | SPORT6 |
| | | including two prima | | | |
| L0641 | NCI_CGAP_Co17 | juvenile granulosa | colon | | pCMV- |
| | | tumor | | | SPORT6 |
| L0643 | NCI_CGAP_Co19 | moderately | colon | | pCMV- |
| | | differentiated | | | SPORT6 |
| L0644 | NCI_CGAP_Co20 | adenocarcinoma | | | 0.01 |
| £0044 | NCI_CGAF_C020 | moderately differentiated | colon | | pCMV- SPORT6 |
| | | adenocarcinoma | | | SPURIO |
| L0645 | NCI_CGAP_Co21 | moderately | colon | | pCMV- |
| ļ | | differentiated | | | SPORT6 |
| | | adenocarcinoma | | | |
| L0646 | NCI_CGAP_Co14 | moderately- | colon | | pCMV- |
| | | differentiated | ĺ | | SPORT6 |
| 1.0645 | NO. COAR G | adenocarcinoma | | | |
| L0647 | NCI_CGAP_Sar4 | five pooled | connective | | pCMV- |
| | | sarcomas, including myxoid liposarcoma | tissue | | SPORT6 |
| L0648 | NCI_CGAP_Eso2 | squamous cell | esophagus | | pCMV- |
| 20010 | TTOI_COM _LDO2 | carcinoma | Csopnagus | | SPORT6 |
| L0649 | NCI_CGAP_GUI | 2 pooled high-grade | genitourinary | | pCMV- |
| | | transitional cell | tract | | SPORT6 |
| | | tumors | | | |
| L0651 | NCI_CGAP_Kid8 | renal cell tumor | kidney | | pCMV- |
| | | | | | SPORT6 |
| L0652 | NCI_CGAP_Lu27 | four pooled poorly- | lung | | pCMV- |
| | | differentiated | | | SPORT6 |
| L0653 | NCI_CGAP_Lu28 | adenocarcinomas | 1,,,,,, | | -CMV |
| T0033 | INCI_CGAP_LUZ8 | two pooled squamous cell | lung | | pCMV- SPORT6 |
| | | carcinomas | | | SLOKIO |
| L0654 | NCI_CGAP_Lu31 | - Can Official | lung, cell line | | pCMV- |
| | , <u> </u> | | | | SPORT6 |
| L0655 | NCI_CGAP_Lym12 | lymphoma, | lymph node | | pCMV- |
| | • | follicular mixed | • • | | SPORT6 |
| | | small and large cell | | | |
| L0656 | NCI_CGAP_Ov38 | normal epithelium | ovary | | pCMV- |

| L0657 | NCI_CGAP_Ov23 | tumor, 5 pooled (see | <u> </u> | _ | SPORT6 |
|----------------|--|-------------------------------------|-------------|--------------|----------------------|
| L0057 | Nei_cdAi _0v23 | description) | ovary | | pCMV- SPORT6 |
| L0658 | NCI_CGAP_Ov35 | tumor, 5 pooled (see | ovary | | pCMV- |
| | | description) | | | SPORT6 |
| L0659 | NCI_CGAP_Pan1 | adenocarcinoma | pancreas | | pCMV- |
| L0661 | NCL CCAD Mails | | | | SPORT6 |
| L0001 | NCI_CGAP_Mel15 | malignant melanoma, | skin | | pCMV- |
| | | metastatic to lymph | | | SPORT6 |
| | | node | | | |
| L0662 | NCI_CGAP_Gas4 | poorly differentiated | stomach | | pCMV- |
| | | adenocarcinoma | | | SPORT6 |
| L0663 | NCI_CGAP_Ut2 | with signet r | | | |
| 10003 | NCI_CGAF_UI2 | moderately- differentiated | uterus | | pCMV- |
| | | endometrial | | | SPORT6 |
| | | adenocarcino | | 1 | |
| L0664 | NCI_CGAP_Ut3 | poorly-differentiated | uterus | | pCMV- |
| | | endometrial | | | SPORT6 |
| L0665 | NCI_CGAP_Ut4 | adenocarcinoma, | | | |
| 1 10003 | NCI_CGAP_U(4 | serous papillary carcinoma, high | uterus | | pCMV- |
| | | grade, 2 pooled t | | | SPORT6 |
| L0666 | NCI_CGAP_Ut1 | well-differentiated | uterus | | pCMV- |
| | | endometrial | | İ | SPORT6 |
| T.066 | | adenocarcinoma, 7 | | | |
| L0667 | NCI_CGAP_CML1 | myeloid cells, 18 | whole blood | | pCMV- |
| | | pooled CML cases, BCR/ABL rearra | | | SPORT6 |
| L0682 | Stanley Frontal NB pool 2 | frontal lobe (see | brain | | pCR2.1- |
| | | description) | Ordin. | | TOPO |
| | | | | | (Invitrogen) |
| L0686 | Stanley Frontal SN pool 2 | frontal lobe (see | brain | | pCR2.1- |
| | | description) , | | İ | ТОРО |
| L0697 | Testis 1 | | | | (Invitrogen) PGEM |
| | | | | | 5zf(+) |
| L0700 | Outward Alu-primed | | *** | *** | pGEM-3Z |
| Y 0715 | hncDNA library | | | | |
| L0717 L0720 | Gessler Wilms tumor PN001-Normal Human | | | | pSPORT1 |
| L0720 | Prostate Prostate | | prostate | | pSport1 |
| L0731 | Soares_pregnant_uterus_ | | uterus | | pT7T3-Pac |
| | NbHPU | | uterus | | p1/13-Pac |
| L0738 | Human colorectal cancer | | | | pT7T3D |
| L0740 | Soares melanocyte | melanocyte | | | pT7T3D |
| | 2NbHM | | |] | (Pharmacia) |
| | | ĺ | | | with a modified |
| | | | | | polylinker |
| L0741 | Soares adult brain | | brain | | pT7T3D |
| i | N2b4HB55Y | ŀ | | | (Pharmacia) |
| | | | | | with a |
| | | 1 | | , | modified |
| L0742 | Soares adult brain | | brain | | polylinker pT7T3D |
| | N2b5HB55Y | , | o.um | | (Pharmacia) |
| | | | | | with a |
| | | | | | modified |
| L0743 | Soares breast 2NbHBst | | 1 | | polylinker |
| 10,43 | Donies Dieast ZINDFIDSt | 1 | breast | | pT7T3D |
| | | | | | (Pharmacia) |

| L0744 Soares breast 3NbHBst breast p1773D (Pharmacia) with a modified polylinker p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) w | <u> </u> | | T | | | |
|---|----------|--------------------------|-------------------|--------------|-----|-------------|
| L0744 Scares breast \$NbHBst Dreast Dolylinker Dol | | | | | • | with a |
| LO745 Soares retina N2b4HR retina eye PTTT3D (Pharmacia) with a modified polylinker poly | İ | | | | | |
| LO745 Soares retina N2b4HR retina eye polylinker polyli | 1.0744 | Soores breest 2NILIDet | | | | |
| L0745 Soares retina N2b4HR retina eye p1773D (Pharmacia) with a modified polylinker p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1 | L0744 | Soales bleast SNoribst | | breast | | |
| L0745 Soares retina N2b4HR retina eye P1773D (Pharmacia) with a modified polylinker poly | 1 | | | 1 | 1 | |
| L0745 Soares retina N2b4HR retina eye polylinker pfT73D (Pharmacia) with a modified polylinker polylinker polylinker profiles a modified polylinker pfT73D (Pharmacia) with a modified pfT73D (Pharmacia) with a modified pfT73D (Pharmacia) with a modified | | | | | | |
| L0746 Soares retina N2b4HR retina eye p1773D (Pharmacia) with a modified polylinker p1773D (Pharmacia) with a modified p0lylinker p1773D (Pharmacia) with a modified p1773D (Pharmacia) with a modified p1773D | | | | | | |
| L0746 Soares retina N2b5HR retina eye printage polylinker pfTri3D (Pharmacia) with a modified pfTri3D (Pharmacia) with a modified pfTri3D (Pharmacia) with a modified | 1.0745 | Soores rating N2h4HD | · · · | | | |
| L0746 Soares retina N2b5HR retina eye pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified pNT3D (Pharmacia) with a modified pNT3D (| 1 20743 | Soares fettila N204AR | retina | eye | | |
| L0746 Soares retina N2b5HR retina eye | Í | | | | | |
| L0746 Soares retina N2b5HR retina eye p7773D (Pharmacia) with a modified polylinker p1773D (Pharmacia) with a modified p0lylinker p1773D (Pharmacia) with a modi | | | | | | |
| L0747 Soares_fetal_heart_NbHH 19W heart p17173D (Pharmacia) with a modified polylinker p17713D (Pharmacia) with a modified p1091 made m0dified p1091 made m0dified p1091 made m0dified p1091 made m0dified m0d | | | • | | | |
| L0747 Soares_fetal_heart_NbHH 19W heart | 1.0746 | Soores rating N255LID | | | | |
| L0747 Soares_fetal_heart_NbHH 19W heart p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified modified polylinker p17T3D (Pharmacia) with a modified mo | L0740 | Soares fettila N2b3rik | retina | eye | | |
| L0747 Soares_fetal_heart_NbHH 19W heart P17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified modified modified modified modified modified modified modified modified modified modified | 1 | | | | | |
| L0747 Soares_fetal_heart_NbHH 19W heart plotylinker plotylinker plotylinker nodified polylinker plotylinker | | | | | |
| LO748 Soares fetal_heart_NbHH Soares fetal_heart_NbHH Soares fetal_liver spleen Liver and spleen PTT3D | | | • | | | |
| LO748 Soares fetal liver spleen Liver and Spleen PTT3D (Pharmacia) with a modified polylinker LO749 Soares_fetal_liver_spleen _ INFLS_S1 Liver and Spleen PTT3D (Pharmacia) with a modified polylinker PTT3D (Pharmacia) with a modified polylinker LO750 Soares_fetal_lung_NbHL1 lung PTT3D (Pharmacia) with a modified polylinker NbHOT Ovary PTT3D (Pharmacia) with a modified polylinker PTT3D (Pharmacia) with a modified polylinker PTT3D (Pharmacia) with a modified PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT3D (PTT | 1.0747 | Connection to the NUTYER | | | | |
| L0748 Soares fetal liver spleen 1NFLS | L0/4/ | | | heart | | |
| Liver and Spleen Liver and Spleen Liver and Spleen Liver and Spleen Liver and Spleen Liver and Spleen Liver and Spleen Liver and Spleen Liver and Spleen Department of the polylinker optimized polylinker optimiz | | 1944 | | | | |
| L0748 Soares fetal liver spleen lNFLS Liver and Spleen Spleen lNFLS Soares fetal liver spleen lNFLS Liver and Spleen Spleen Spleen lnFLS_S1 Liver and Spleen Spleen PDT713D (Pharmacia) with a modified polylinker pT713D (Pharmacia) with a modified polylinker pNHL1 swith a modified polylinker pNHD1 swith a modified polylinker pNHD1 swith a modified polylinker pNHD1 swith a modified polylinker pNHD1 swith a modified polylinker pNHD1 swith a modified polylinker pNHD2 soares parathyroid_tumor NbHD4 profit polylinker p | | } | 1 | | 1 | |
| Liver and Spleen INFLS Soares fetal liver spleen INFLS Soares_fetal_liver_spleen Liver and Spleen PT73D (Pharmacia) with a modified polylinker | ł | | | | | modified |
| INFLS Spleen PATTAGIA With a modified polylinker p1773D (Pharmacia) with a modified modifi | 1.0749 | Constitution 1 | | | | |
| L0750 Soares_fetal_liver_spleenINFLS_S1 | LU/48 | | | | 1 | pT7T3D |
| L0749 Soares_fetal_liver_spleenINFLS_S1 | | INFLS | | Spleen | | (Pharmacia) |
| L0749 Soares_fetal_liver_spleen | İ | | | | 1 | with a |
| Liver and Spleen Liver and Spleen PTT3D (Pharmacia) with a modified polylinker polyl | | | | | 1 | modified |
| L0750 Soares fetal_lung_NbHL1 PG Soares placenta Nb2HP Soares placenta Spleen Spleen Spleen (Pharmacia) with a modified polylinker placenta Nb2HP Soares placenta_Sto9we ks_2NbHP8to9W Soares_placenta_Sto9we ks_2NbHP8to9W Soares_placenta_Sto9we ks_2NbHP8to9W Soares_placenta_Sto9were placenta NbHP8to9W Soares_placenta_Sto9were placenta NbHP8to9W Soares_placenta_Sto9were placenta pland pland placenta | 1.0740 | | | | | polylinker |
| L0750 Soares_fetal_lung_NbHL1 | L0749 | | | | | pT7T3D |
| L0750 Soares_fetal_lung_NbHL1 | ļ | _INFLS_S1 | | Spleen | | (Pharmacia) |
| L0750 Soares_fetal_lung_NbHL1 9W lung pT7T3D (Pharmacia) with a modified polylinker pOlylinker pOlylinker poly | i | | | | | with a |
| LU751 Soares_letal_lung_NbHL1 9W lung pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker polylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a modified pOlylinker pT7T3D (Pharmacia) with a | | | | | | modified |
| 9W LO751 Soares ovary tumor NbHOT ovarian tumor ovary pT7T3D (Pharmacia) with a modified polylinker LO752 Soares_parathyroid_tumorNbHPA pG LO753 Soares_pineal_gland_N3H PG LO754 Soares_placenta_Nb2HP placenta LO755 Soares_placenta_Sto9wee ks_2NbHP8to9W POTT3D (Pharmacia) with a modified polylinker placenta p | 1.0750 | | | | | polylinker |
| L0751 Soares ovary tumor NbHOT ovarian tumor ovary pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pOlylinker pOlylinker pOlylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified | L0750 | | | lung | | pT7T3D |
| L0751 Soares ovary tumor NbHOT ovarian tumor ovary pT7T3D (Pharmacia) with a modified polylinker L0752 Soares_parathyroid_tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA pineal gland p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified | ł | 9W | | | | (Pharmacia) |
| L0751 Soares ovary tumor NbHOT | 1 | 1 | | 1 | | with a |
| L0751 Soares ovary tumor NbHOT ovarian tumor ovary pT7T3D (Pharmacia) with a modified polylinker L0752 Soares_parathyroid_tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid gland p17T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified polylinker p17T3D (Pharmacia) with a modified | | | | 1 | | |
| NbHOT NbHOT NbHOT NbHOT NbHOT NbHPA Parathyroid tumor NbHPA Parathyroid tumor NbHPA Parathyroid tumor NbHPA Parathyroid tumor NbHPA Parathyroid tumor NbHPA Parathyroid tumor NbHPA Parathyroid tumor NbHPA Parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid gland parathyroid parathyroid gland parathyroid parathyroid parathyroid parathyroid gland parathyroid par | T 0751 | | | | | |
| L0752 Soares_parathyroid_tumorNbHPA | LU/51 | Soares ovary tumor | ovarian tumor | ovary | | pT7T3D |
| L0752 Soares_parathyroid_tumorNbHPA | İ | NBHOT | | | | |
| L0752 Soares_parathyroid_tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid tumor _NbHPA parathyroid gland polylinker _NbHPA polylinker pineal gland polylinker pineal gland polylinker pineal gland polylinker pineal placenta pineal polylinker pineal placenta pineal pineal polylinker pineal placenta pineal pine | | | | | | with a |
| L0752 Soares_parathyroid_tumor | | | | | .] | modified |
| NbHPA gland (Pharmacia) with a modified polylinker PG pineal gland pineal gland (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker placenta Nb2HP placenta pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified with a modified with a modified polylinker pT7T3D (Pharmacia) with a modified | 1.0750 | | | | | polylinker |
| L0753 Soares_pineal_gland_N3H PG pineal gland pT7T3D (Pharmacia) with a modified polylinker L0754 Soares placenta Nb2HP placenta p17T3D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta p17T3D (Pharmacia) with a modified with a modified polylinker | LU/52 | Soares_parathyroid_tumor | parathyroid tumor | parathyroid | 1 | pT7T3D |
| L0753 Soares_pineal_gland_N3H pineal gland pineal gland pT7T3D (Pharmacia) with a modified polylinker L0754 Soares placenta Nb2HP placenta placenta pT7T3D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pT7T3D (Pharmacia) with a modified with a modified polylinker | 1 | _NbHPA | | gland | 1 | (Pharmacia) |
| L0753 Soares_pineal_gland_N3H PG pineal gland pineal gland pT7T3D (Pharmacia) with a modified polylinker L0754 Soares placenta Nb2HP placenta pT7T3D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta pT7T3D (Pharmacia) with a modified with a modified polylinker | | | | | | |
| L0753 Soares_pineal_gland_N3H PG pineal gland pineal gland pT7T3D (Pharmacia) with a modified polylinker L0754 Soares placenta Nb2HP placenta pT7T3D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta pT7T3D (Pharmacia) with a modified with a modified polylinker | 1 | | | | | modified |
| PG PG PG PG PG PG PG Pharmacia) With a modified polylinker PTT3D (Pharmacia) With a modified With a modified PTT3D (Pharmacia) With a modified | 1.0752 | | | | | |
| L0754 Soares placenta Nb2HP placenta placenta placenta placenta pr773D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pr773D (Pharmacia) with a modified with a modified polylinker pr773D (Pharmacia) with a modified | LU/53 | | | pineal gland | | |
| L0754 Soares placenta Nb2HP placenta placenta pT7T3D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pT7T3D (Pharmacia) with a modified with a modified polylinker pT7T3D (Pharmacia) with a modified | | PG | | | 1 | |
| L0754 Soares placenta Nb2HP placenta placenta pplylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pplylinker ppr7T3D (Pharmacia) with a modified polylinker ppr7T3D (Pharmacia) with a modified | | | | | | |
| L0754 Soares placenta Nb2HP placenta placenta pT7T3D (Pharmacia) with a modified polylinker L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta pT7T3D (Pharmacia) with a modified with a modified polylinker pT7T3D (Pharmacia) with a modified | Į. | } | | | 1 1 | |
| L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pT7T3D (Pharmacia) with a modified polylinker pT7T3D (Pharmacia) with a modified | 1.0754 | Constant | | | | |
| L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pT7T3D (Pharmacia) with a modified polylinker | LU/54 | Soares placenta Nb2HP | | placenta | | |
| L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta placenta pT7T3D (Pharmacia) with a modified | | | | | | |
| L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta pplacenta pplacenta ppr7T3D (Pharmacia) with a modified |] | | | | | |
| L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta ppT7T3D (Pharmacia) with a modified | | | | | | |
| L0755 Soares_placenta_8to9wee ks_2NbHP8to9W placenta ppT7T3D (Pharmacia) with a modified | 1077 | | | | | |
| ks_2NbHP8to9W (Pharmacia) with a modified | L0755 | | | placenta | | pT7T3D |
| with a modified |] | ks_2NbHP8to9W | | 1 | | |
| | | · | | | | with a |
| polylinker | | | | | | |
| | | <u></u> | | | | polylinker |

| | | · | | | | |
|----------|---------------------------|----------------------|--------|---|----------------|---------------------------|
| L0756 | Soares_multiple_sclerosis | multiple sclerosis | | | | pT7T3D |
| İ | _2NbHMSP | lesions | | | | (Pharmacia) |
| | | j | | | | with a |
| - | | | | | | modified |
| 1 | | | | | | polylinker |
| 1 | | | | İ | | V_TYPE |
| L0757 | Soares_senescent_fibrobla | senescent fibroblast | | | | pT7T3D |
| | sts_NbHSF | | | | ļ | (Pharmacia) |
| | | | | ł | ĺ | with a |
| | | | | | ļ | |
| 1 | | | | j | | modified |
| | | | | | İ | polylinker |
| L0758 | Soares_testis_NHT | | | | ļ | V_TYPE |
| L0/36 | Soales_lestis_NH1 | | | | | pT7T3D-Pac |
| | | ì | | | | (Pharmacia) |
| | | | | | | with a |
| 1 | | | | | | modified |
| | | | | | | polylinker |
| L0759 | Soares_total_fetus_Nb2H | | | | | pT7T3D-Pac |
| | F8_9w | | | j | | (Pharmacia) |
| | | | | | | with a |
| | | 1 | | | | modified |
| | | | | | | polylinker |
| L0761 | NCI_CGAP_CLL1 | B-cell, chronic | | | | pT7T3D-Pac |
| | | lymphotic leukemia | | | 1 | (Pharmacia) |
| | 1 | | | | | with a |
| | | | 1 | | | modified |
| | | [| | | | |
| L0762 | NCI_CGAP_Br1.1 | breast | | | | polylinker |
| 20702 | Nei_eg/ii _Bi1.i | bleast | | | 1 | pT7T3D-Pac |
| | | | | | | (Pharmacia) |
| | | | | | | with a |
| | | | | | | modified |
| 1.0762 | NOT COAD DO | | | | | polylinker |
| L0763 | NCI_CGAP_Br2 | breast | | | ļ | pT7T3D-Pac |
| | | | | į | | (Pharmacia) |
| | | | | | | with a |
| | | i | | | | modified |
| | | | | | | polylinker |
| L0764 | NCI_CGAP_Co3 | colon | | | | pT7T3D-Pac |
| | | | • | 1 | | (Pharmacia) |
| | | | | | 1 | with a |
| | | | | | | modified |
| | | | | | | polylinker |
| L0766 | NCI_CGAP_GCB1 | germinal center B | | | | pT7T3D-Pac |
| | | cell | | | | (Pharmacia) |
| | | · | | | | with a |
| | | | | | | modified |
| | | | | 1 | | polylinker |
| L0767 | NCI_CGAP_GC3 | pooled germ cell | | | - | |
| | | tumors | | | 1 | pT7T3D-Pac |
| | | rantors | | 1 | | (Pharmacia) |
| | | | | 1 | | with a |
| | | | | | | modified |
| L0768 | NCI_CGAP_GC4 | noolod 11 | | | - | polylinker |
| LU/00 | NCI_COAF_GC4 | pooled germ cell | | 1 | | pT7T3D-Pac |
| | | tumors | | 1 | | (Pharmacia) |
| | | | | | | with a |
| | | | | i | | modified |
| | | | | | | polylinker |
| L0769 | NCI_CGAP_Brn25 | anaplastic | brain | | | pT7T3D-Pac |
| | | oligodendroglioma | | | | (Pharmacia) |
| | l | ' | | | | with a |
| i | | | | | | modified |
| | | | | 1 | | polylinker |
| L0770 | NCI_CGAP_Brn23 | glioblastoma | brain | 1 | | |
| L01/0 | 1101_CG/11 _B11123 | Enoplasiona | Ulalli | | | י סוייים איינונין/ום |
| LOTTO | · | (pooled) | Oram | 1 | | pT7T3D-Pac (Pharmacia) |

| | | T | | | |
|---------|------------------------|-----------------------------|-------------|--|-------------|
| | | | | | with a |
| | | | | | modified |
| | | | | | polylinker |
| L0771 | NCI_CGAP_Co8 | adenocarcinoma | colon | | pT7T3D-Pac |
| | | | | | (Pharmacia) |
| | | | | | with a |
| 1 | | | | | modified |
| | | | | 1 | polylinker |
| L0772 | NCI_CGAP_Co10 | colon tumor RER+ | colon | | pT7T3D-Pac |
| | | | | | (Pharmacia) |
| | | 1 | | | with a |
| | | | | | modified |
| 1 | | | | | polylinker |
| L0773 | NCI_CGAP_Co9 | colon tumor RER+ | colon | | pT7T3D-Pac |
| | | Total turnor representation | Colon | | (Pharmacia) |
| 1 | <u> </u> | 1 | | | with a |
| | | | | | modified |
| | | | | | |
| L0774 | NCI_CGAP_Kid3 | | kidney | | polylinker |
| 1 20774 | Nei_edAi_Rid5 | | Kidney | | pT7T3D-Pac |
| 1 | | i | | | (Pharmacia) |
| | | | 1 | | with a |
| 1 | | | 1 | | modified |
| 1.0775 | NOT COAD YELD | | ļ | | polylinker |
| L0775 | NCI_CGAP_Kid5 | 2 pooled tumors | kidney | | pT7T3D-Pac |
| | | (clear cell type) | | | (Pharmacia) |
| | | | | | with a |
| 1 | | | | | modified |
| | | | | | polylinker |
| L0776 | NCI_CGAP_Lu5 | carcinoid | lung | | pT7T3D-Pac |
| | | | | | (Pharmacia) |
| i | | 4 | | | with a |
| | | | | | modified |
| | | | | | polylinker |
| L0777 | Soares_NhHMPu_S1 | Pooled human | mixed (see | | pT7T3D-Pac |
| - | _ | melanocyte, fetal | below) | | (Pharmacia) |
| | | heart, and pregnant | | 1 | with a |
| | · |] | | 1 | modified |
| 1 | | ļ |] | 1 | polylinker |
| L0779 | Soares_NFL_T_GBC_S1 | | pooled | | pT7T3D-Pac |
| 1 | 000.00_1112_12020_01 | | pooled | | (Pharmacia) |
| | | | | İ | with a |
| | | | | | modified |
| 1 | | | | | |
| L0780 | Soares_NSF_F8_9W_OT | | mac1-3 | | polylinker |
| L0760 | _PA_P_S1 | , | pooled | | pT7T3D-Pac |
| | | , | | | (Pharmacia) |
| | | | | | with a |
| 1 | (| | 1 | | modified |
| T 0702 | NOI COAD DOG | | | | polylinker |
| L0783 | NCI_CGAP_Pr22 | normal prostate | prostate | | pT7T3D-Pac |
| | | | | | (Pharmacia) |
| l | · . | | | | with a |
| | | | | | modified |
| L | 3.03 | | | | polylinker |
| L0784 | NCI_CGAP_Lei2 | leiomyosarcoma | soft tissue | | pT7T3D-Pac |
| [| | | | l İ | (Pharmacia) |
| | | | • | | with a |
| | | | | | modified |
| | | | | | polylinker |
| L0785 | Barstead spleen HPLRB2 | | spleen | | pT7T3D-Pac |
| | · | | 1 | | (Pharmacia) |
| | | | | | with a |
| | - | | | | modified |
| | | | | | polylinker |
| | <u> </u> | | | L | Poryminer |

| 1.0706 | 10 MILLED | | | | |
|----------|-----------------|--------------------|---------------------------------------|--|-------------|
| L0786 | Soares_NbHFB | | whole brain | | pT7T3D-Pac |
| | ļ | | | | (Pharmacia) |
| İ | Ì | | | | with a |
| | | | | | modified |
| L | | | | | polylinker |
| L0787 | NCI_CGAP_Sub1 | | | | pT7T3D-Pac |
| İ | 1 | | 1 | 1 | (Pharmacia) |
| | | | | 1 | with a |
| | 1 | | | | modified |
| | | | | 1 | polylinker |
| L0788 | NCI_CGAP_Sub2 | | · · · · · · · · · · · · · · · · · · · | | |
| | 11012001112002 | | | 1 | pT7T3D-Pac |
| | [| | | | (Pharmacia) |
| | | | | | with a |
| 1 | i | | | | modified |
| L0789 | NCI_CGAP_Sub3 | | | <u> </u> | polylinker |
| L0789 | NCI_CGAP_Sub3 | İ | | 1 | pT7T3D-Pac |
| 1 | Ì | | | | (Pharmacia) |
| 1 | | | | | with a |
| | | | | | modified |
| | | | | | polylinker |
| L0790 | NCI_CGAP_Sub4 | | | | pT7T3D-Pac |
| 1 | | 1 | |] | (Pharmacia) |
|] | į | | |] | with a |
| | | | | 1 | modified |
| | | ļ | | 1 1 | polylinker |
| L0791 | NCI_CGAP_Sub5 | | | | |
| | | | | 1 | pT7T3D-Pac |
| | | | | 1 | (Pharmacia) |
| | ļ | Į | | | with a |
| 1 | | | | | modified |
| L0792 | NCI_CGAP_Sub6 | | | | polylinker |
| L0792 | NCI_CGAP_Subb | | | | pT7T3D-Pac |
| | | | | | (Pharmacia) |
| 1 | 1 | | ĺ | 1 | with a |
| | | | | | modified |
| | *** | | | | polylinker |
| L0794 | NCI_CGAP_GC6 | pooled germ cell | | | pT7T3D-Pac |
| ſ | | tumors | | | (Pharmacia) |
| | ļ | | | | with a |
| | İ | | | | modified |
| ł | | | | | polylinker |
| L0796 | NCI_CGAP_Brn50 | medulloblastoma | brain | | pT7T3D-Pac |
| | | | 01444 | | (Pharmacia) |
| l | | | | | with a |
| | | | | | |
| | ŀ | | | } | modified |
| L0800 | NCI_CGAP_Co16 | colon tumor, RER+ | colon | | polylinker |
| 20000 | 1.01_00/11_0010 | colon tullor, KER+ | COLOII | | pT7T3D-Pac |
| 1 | 1 | 1 | | 1 | (Pharmacia) |
| | | | | | with a |
| | | | | | modified |
| T 0000 | NGL GGAE TELL | | | | polylinker |
| L0803 | NCI_CGAP_Kid11 | | kidney | | pT7T3D-Pac |
| 1 | | | | 1 | (Pharmacia) |
| | | | | 1 | with a |
| i | | į į | | 1 | modified |
| | | | | | polylinker |
| L0804 | NCI_CGAP_Kid12 | 2 pooled tumors | kidney | | pT7T3D-Pac |
| | | (clear cell type) | 1 | [| (Pharmacia) |
| | | '' | Ì | İ | with a |
| | | į į | , | 1 | modified |
| | | | | | i i |
| L0805 | NCI_CGAP_Lu24 | carcinoid | lung | | polylinker |
| | O OO. II _Luz- | Carcinoid | inita | | pT7T3D-Pac |
| | | 1 | 1 | 1 | (Pharmacia) |
| | | 1 | ĺ | | with a |
| | | _1 | | | modified |

| | | | | | polylinker |
|-------|--|---|----------|----|---|
| L0806 | NCI_CGAP_Lu19 | squamous cell carcinoma, poorly differentiated (4 | lung | Ŷ. | pT7T3D-Pac (Pharmacia) with a modified |
| | | , | | | polylinker |
| L0809 | NCI_CGAP_Pr28 | | prostate | | pT7T3D-Pac (Pharmacia) with a modified polylinker |
| L2245 | NEM subtracted human fetal kidney cDNA | | | | pUEX1 |
| L2250 | Human cerebral cortex | cerebral cortex | | | |
| L2251 | Human fetal lung | Fetal lung | | | |

TABLE 5

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| 109270 | Hemolytic anemia due to band 3 defect |
|--------|---|
| 109560 | Leukemia/lymphoma, B-cell, 3 |
| 109690 | Asthma, nocturnal, susceptibility to |
| 109690 | Obesity, susceptibility to |
| 109700 | Hemodialysis-related amyloidosis |
| 110700 | Vivax malaria, susceptibility to |
| 113705 | Ovarian cancer |
| 113705 | Breast cancer-1 |
| 113721 | Breast cancer |
| 113900 | Heart block, progressive familial, type I |
| 114208 | Malignant hyperthermia susceptibility 5, 601887 |
| 114208 | Hypokalemic periodic paralysis, 170400 |
| 114350 | Leukemia, acute myeloid |
| 114835 | Monocyte carboxyesterase deficiency |
| 115470 | Cat eye syndrome |
| 116800 | Cataract, Marner type |
| 116860 | Cavernous angiomatous malformations |
| 117700 | [Hypoceruloplasminemia, hereditary] |
| 117700 | Hemosiderosis, systemic, due to aceruloplasminemia |
| 118210 | Charcot-Marie-Tooth neuropathy-2A |
| 118485 | Polycystic ovary syndrome with hyperandrogenemia |
| 119300 | van der Woude syndrome |
| 120110 | Metaphyseal chondrodysplasia, Schmid type |
| 120215 | Ehlers-Danlos syndrome, type I, 130000 |
| 120215 | Ehlers-Danlos syndrome, type II, 130010 |
| 120220 | Bethlem myopathy, 158810 |
| 120240 | Bethlem myopathy, 158810 |
| 120260 | Epiphyseal dysplasia, multiple, type 2, 600204 |
| 120280 | Stickler syndrome, type III |
| 120280 | Marshall syndrome, 154780 |
| 120290 | OSMED syndrome, 215150 |
| 120290 | Stickler syndrome, type II, 184840 |
| 120435 | Muir-Torre syndrome, 158320 |
| 120435 | Colorectal cancer, hereditary, nonpolyposis, type 1 Ovarian |
| | cancer |
| 120550 | C1q deficiency, type A |
| 120570 | C1q deficiency, type B |
| 120575 | C1q deficiency, type C |
| 120620 | SLE susceptibility |
| 120620 | CR1 deficiency |
| 120700 | C3 deficiency |
| 120810 | C4 deficiency |
| 120820 | C4 deficiency |
| 120900 | C5 deficiency |
| 120920 | Measles, susceptibility to |
| 120940 | C9 deficiency |
| | |

| 120960 C8 deficiency, type II | 120950 | C8 deficiency, type I |
|--|--------|--|
| 121011 Deafness, autosomal dominant 3, 601544 121011 Deafness, autosomal recessive 1, 220290 121014 Heterotaxia, visceroatrial, autosomal recessive 1 121360 Myeloid leukemia, acute, M4Eo subtype 121700 Congenital hereditary endothelial dystrophy of cornea 121800 Corneal dystrophy, crystalline, Schnyder 122000 Corneal dystrophy, posterior polymorphous 122500 [Transcortin deficiency] 122560 ACTH deficiency, 201400 122720 Nicotine addiction, protection from 122720 Nicotine addiction, protection from 122720 Coumarin resistance, 122700 123000 Craniometaphyscal dysplasia 123101 Craniosynostosis, type 2 123270 [Creatine kinase, brain type, ectopic expression of] 123580 Cataract, congenital, autosomal dominant 123620 Cataract, cerulean, type 2, 601547 123829 Melanoma 123940 White sponge nevus, 193900 124030 Parkinsonism, susceptibility to 124030 Debrisoquine sensitivity 124200 Darier disease (keratosis follicularis) 125490 Dentinogenesis imperfecta-1 125852 Insulin-dependent diabetes mellitus-2 126060 Anemia, megaloblastic, due to DHFR deficiency 126391 DNA ligase I deficiency 126452 Autonomic nervous system dysfunction 12659 Chloride diarrhea, congenital, Finnish type, 214700 12650 Colon cancer 129490 Ectodermal dysplasia-3, anhidrotic 129900 EEC syndrome-1 130410 Glutaricaciduria, type IIB 130500 Elliptocytosis-1 | | |
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| 131195 Hereditary hemorrhagic telangiectasia-1, 187300 | | |
| 131210 Atherosclerosis, susceptibility to | | |
| 131400 Eosinophilia, familial | | |
| 132700 Cylindromatosis | 132700 | Cylindromatosis |
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| 133171 [Erythrocytosis, familial], 133100 | 133171 | [Erythrocytosis, familial], 133100 |
| 133200 Erythrokeratodermia variabilis | | |

| 133780 | Vitreoretinopathy, exudative, familial |
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| 134370 | Factor H deficiency |
| 134370 | Hemolytic-uremic syndrome, 235400 |
| 134370 | Membroproliferative glomerulonephritis |
| 134570 | Factor XIIIA deficiency |
| 134580 | Factor XIIIB deficiency |
| 134790 | Hyperferritinemia-cataract syndrome, 600886 |
| 135300 | Fibromatosis, gingival |
| 135940 | Ichthyosis vulgaris, 146700 |
| 136132 | [Fish-odor syndrome], 602079 |
| 136435 | Ovarian dysgenesis, hypergonadotropic, with normal karyotype, 233300 |
| 136550 | Macular dystrophy, North Carolina type |
| 136836 | Fucosyltransferase-6 deficiency |
| 137600 | Iridogoniodysgenesis syndrome |
| 138079 | Hyperinsulinism, familial, 602485 |
| 138079 | MODY, type 2, 125851 |
| 138140 | Glucose transport defect, blood-brain barrier |
| 138190 | Diabetes mellitus, noninsulin-dependent |
| 138491 | Startle disease, autosomal recessive |
| 138491 | Startle disease/hyperekplexia, autosomal dominant, 149400 |
| 138491 | Hyperekplexia and spastic paraparesis |
| 138570 | Non-insulin dependent diabetes mellitus, susceptibility to |
| 138971 | Kostmann neutropenia, 202700 |
| 138981 | Pulmonary alveolar proteinosis, 265120 |
| 139190 | Gigantism due to GHRF hypersecretion |
| 139190 | Isolated growth hormone deficiency due to defect in GHRF |
| 139191 | Growth hormone deficient dwarfism |
| 139330 | Night blindness, congenital stationary |
| 139350 | Epidermolytic hyperkeratosis, 113800 |
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| 139360 | Pituitary ACTH-secreting adenoma |
| 140100 | [Anhaptoglobinemia] |
| 140100 | [Hypohaptogloginemia] |
| 141900 | Methemoglobinemias, beta- |
| 141900 | Sickle cell anemia |
| 141900 | Thalassemias, beta- |
| 141900 | Erythremias, beta- |
| 141900 | HPFH, deletion type |
| 141900 | Heinz body anemias, beta- |
| 142000 | Thalassemia due to Hb Lepore |
| 142000 | Thalassemia, delta- |
| 142200 | HPFH, nondeletion type A |
| 142250 | HPFH, nondeletion type G |
| 142270 | Hereditary persistence of fetal hemoglobin |
| 142360 | Thrombophilia due to heparin cofactor II deficiency |

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| Epidermolysis bullosa, generalized atrophic benign, 226650 | 150310 | Epidermolysis bullosa, Herlitz junctional type, 226700 |
| | | Epidermolysis bullosa, generalized atrophic benign, 226650 |
| | 151385 | |

| 151410 | Leukemia, chronic myeloid |
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| 151440 | Leukemia, T-cell acute lymphoblastoid |
| 151670 | Hepatic lipase deficiency |
| 152200 | Coronary artery disease, susceptibility to |
| 152445 | Vohwinkel syndrome, 124500 |
| 152445 | Erythrokeratoderma, progressive symmetric, 602036 |
| 152760 | Hypogonadotropic hypogonadism due to GNRH deficiency, 227200 |
| 152780 | Hypogonadism, hypergonadotropic |
| 152780 | Male pseudohermaphroditism due to defective LH |
| 152790 | Precocious puberty, male, 176410 |
| 152790 | Leydig cell hypoplasia |
| 153455 | Cutis laxa, recessive, type I, 219100 |
| 153880 | Macular dystrophy, dominant cystoid |
| 154275 | Malignant hyperthermia susceptibility 2 |
| 154276 | Malignant hyperthermia susceptibility 3 |
| 154500 | Treacher Collins mandibulofacial dysostosis |
| 154550 | Carbohydrate-deficient glycoprotein syndrome, type Ib, 602579 |
| 156225 | Muscular dystrophy, congenital merosin-deficient |
| 157170 | Holoprosencephaly-2 |
| 157640 | PEO with mitochondrial DNA deletions, type 1 |
| 159000 | Muscular dystrophy, limb-girdle, type 1A |
| 159001 | Muscular dystrophy, limb-girdle, type 1B |
| 159440 | Charcot-Marie-Tooth neuropathy-1B, 118200 |
| 159440 | Dejerine-Sottas disease, myelin P-related, 145900 |
| 159440 | Hypomyelination, congenital |
| 160781 | Cardiomyopathy, hypertrophic, mid-left ventricular chamber type |
| 160900 | Myotonic dystrophy |
| 162200 | Neurofibromatosis, type 1 |
| 162200 | Watson syndrome, 193520 |
| 164040 | Leukemia, acute promyelocytic, NPM/RARA type |
| 164200 | Oculodentodigital dysplasia |
| 164200 | Syndactyly, type III, 186100 |
| 164500 | Spinocerebellar ataxia-7 |
| 164731 | Ovarian carcinoma, 167000 |
| 164860 | Renal cell carcinoma, papillary, familial and sporadic |
| 164953 | Liposarcoma |
| 165215 | 3q21q26 syndrome |
| 165500 | Optic atrophy 1 |
| 166600 | Osteopetrosis, AD, type II |
| 166800 | Otosclerosis Otosclerosis |
| 167000 | Ovarian cancer, serous |
| 167250 | Paget disease of bone |
| 167409 | Optic nerve coloboma with renal disease, 120330 |
| 167415 | Hypothyroidism, congenital, due to thyroid dysgenesis or |
| 10/115 | hypoplasia die to thyroid dysgenesis or |
| | 1 17 Poptasta |

| 168360 | Downwardsting |
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| 168468 | Paraneoplastic sensory neuropathy |
| 168610 | Metaphyseal chondrodysplasia, Murk Jansen type, 156400 |
| 169600 | Parkinsonism-dementia with pallidopontonigral degeneration |
| 170261 | Hailey-Hailey disease |
| | Bare lymphocyte syndrome, type I, due to TAP2 deficiency |
| 170650 | Periodontitis, juvenile |
| 170995 | Zellweger syndrome-2 |
| 171190 | Hypertension, essential, 145500 |
| 171760 | Hypophosphatasia, adult, 146300 |
| 171760 | Hypophosphatasia, infantile, 241500 |
| 171860 | Hemolytic anemia due to phosphofructokinase deficiency |
| 172400 | Hemolytic anemia due to glucosephosphate isomerase deficiency |
| 172400 | Hydrops fetalis, one form |
| 172411 | Colorectal cancer, resistance to |
| 172471 | Glycogenosis, hepatic, autosomal |
| 172490 | Phosphorylase kinase deficiency of liver and muscle, 261750 |
| 173350 | Plasminogen Tochigi disease |
| 173350 | Plasminogen deficiency, types I and II |
| 173350 | Thrombophilia, dysplasminogenemic |
| 173360 | Thrombophilia due to excessive plasminogen activator inhibitor |
| 173360 | Hemorrhagic diathesis due to PAI1 deficiency |
| 173610 | Platelet alpha/delta storage pool deficiency |
| 173850 | Polio, susceptibility to |
| 174000 | Medullary cystic kidney disease, AD |
| 174900 | Polyposis, juvenile intestinal |
| 176000 | Porphyria, acute intermittent |
| 176100 | Porphyria cutanea tarda |
| 176100 | Porphyria, hepatoerythropoietic |
| 176705 | Breast cancer, sporadic |
| 176730 | Diabetes mellitus, rare form |
| 176730 | Hyperproinsulinemia, familial |
| 176730 | MODY, one form |
| 176830 | Obesity, adrenal insufficiency, and red hair |
| 176830 | ACTH deficiency |
| 176860 | Purpura fulminans, neonatal |
| 176860 | Thrombophilia due to protein C deficiency |
| 176943 | Apert syndrome, 101200 |
| 176943 | Pfeiffer syndrome, 101600 |
| 176943 | Beare-Stevenson cutis gyrata syndrome, 123790 |
| 176943 | Crouzon craniofacial dysostosis, 123500 |
| 176943 | Jackson-Weiss syndrome, 123150 |
| 176947 | Selective T-cell defect |
| 177070 | Spherocytosis, hereditary, Japanese type |
| 177070 | Hermansky-Pudlak syndrome, 203300 |
| 177900 | Psoriasis susceptibility-1 |
| 178300 | |
| 1/02/// | Ptosis, hereditary congenital, 1 |

| 178640 | Pulmonary alveolar proteinosis, congenital, 265120 |
|--------|--|
| 179450 | Ragweed sensitivity |
| 179755 | Renal cell carcinoma, papillary, 1 |
| 179820 | [Hyperproreninemia] |
| 180020 | Retinal cone dystrophy-1 |
| 180071 | Retinitis pigmentosa, autosomal recessive |
| 180100 | Retinitis pigmentosa-1 |
| 180104 | Retinitis pigmentosa-9 |
| 180105 | Retinitis pigmentosa-10 |
| 180250 | Retinol binding protein, deficiency of |
| 180297 | Anemia, hemolytic, Rh-null, suppressor type, 268150 |
| 180380 | Night blindness, congenital stationery, rhodopsin-related |
| 180380 | Retinitis pigmentosa, autosomal recessive |
| 180380 | Retinitis pigmentosa-4, autosomal dominant |
| 180901 | Malignant hyperthermia susceptibility 1, 145600 |
| 180901 | Central core disease, 117000 |
| 181405 | Scapuloperoneal spinal muscular atrophy, New England type |
| 181430 | Scapuloperoneal syndrome, myopathic type |
| 181460 | Schistosoma mansoni, susceptibility/resistance to |
| 181510 | Schizophrenia |
| 182138 | Anxiety-related personality traits |
| 182280 | Small-cell cancer of lung |
| 182380 | Glucose/galactose malabsorption |
| 182381 | Renal glucosuria, 253100 |
| 182500 | Cataract, congenital |
| 182600 | Spastic paraplegia-3A |
| 182601 | Spastic paraplegia-4 |
| 182860 | Pyropoikilocytosis |
| 182860 | Spherocytosis, recessive |
| 182860 | Elliptocytosis-2 |
| 185000 | Stomatocytosis I |
| 185430 | Atherosclerosis, susceptibility to |
| 185470 | Myopathy due to succinate dehydrogenase deficiency |
| 185800 | Symphalangism, proximal |
| 186580 | Arthrocutaneouveal granulomatosis |
| 186770 | Leukemia, T-cell acute lymphocytic |
| 186780 | CD3, zeta chain, deficiency |
| 186880 | Leukemia/lymphoma, T-cell |
| 186921 | Leukemia, T-cell acute lymphoblastic |
| 186960 | Leukemia/lymphoma, T-cell |
| 187040 | Leukemia-1, T-cell acute lymphoblastic |
| 188040 | Thrombophilia due to thrombomodulin defect |
| 188070 | Bleeding disorder due to defective thromboxane A2 receptor |
| 188400 | Velocardiofacial syndrome, 192430 |
| 188400 | DiGeorge syndrome |
| 188826 | Sorsby fundus dystrophy, 136900 |

| 189800 | Preeclampsia/eclampsia |
|---------|--|
| 189980 | Leukemia, chronic myeloid |
| 190000 | Atransferrinemia |
| 190020 | Bladder cancer, 109800 |
| 190040 | Dermatofibrosarcoma protuberans |
| 190040 | Giant-cell fibroblastoma |
| 190040 | Meningioma, SIS-related |
| 190100 | Geniospasm |
| 190195 | Ichthyosiform erythroderma, congenital, 242100 |
| 190195 | Ichthyosis, lamellar, autosomal recessive, 242300 |
| 190198 | Leukemia, T-cell acute lymphoblastic |
| 190685 | Down syndrome |
| 190900 | Colorblindness, tritan |
| 191030 | Nemaline myopathy-1, 161800 |
| 191044 | Cardiomyopathy, familial hypertrophic |
| 191045 | Cardiomyopathy, familial hypertrophic, 2, 115195 |
| 191290 | Segawa syndrome, recessive |
| 191315 | Insensitivity to pain, congenital, with anhidrosis, 256800 |
| 192090 | Ovarian carcinoma |
| 192090 | Breast cancer, lobular |
| 192090 | Endometrial carcinoma |
| 192090 | Gastric cancer, familial, 137215 |
| 192500 | Jervell and Lange-Nielsen syndrome, 220400 |
| 192500 | Long QT syndrome-1 |
| 194071 | Wilms tumor, type 2 |
| 194071 | Adrenocortical carcinoma, hereditary, 202300 |
| 200150 | Choreoacanthocytosis |
| 200350 | Acetyl-CoA carboxylase deficiency |
| 201470 | Acyl-CoA dehydrogenase, short-chain, deficiency of |
| 201910 | Adrenal hyperplasia, congenital, due to 21-hydroxylase |
| | deficiency |
| 202200 | Glucocorticoid deficiency, due to ACTH unresponsiveness |
| 203310 | Ocular albinism, autosomal recessive |
| 203500 | Alkaptonuria |
| 204500 | Ceroid-lipofuscinosis, neuronal 2, classic late infantile |
| 205900 | Anemia, Diamond-Blackfan |
| 207750 | Hyperlipoproteinemia, type Ib |
| -207800 | Argininemia |
| 208100 | Arthrogryposis multiplex congenita, neurogenic |
| 208250 | Jacobs syndrome |
| 210900 | Bloom syndrome |
| 214400 | Charcot-Marie-Tooth neuropathy-4A |
| 216550 | Cohen syndrome |
| 216900 | Achromatopsia |
| 217000 | C2 deficiency |
| 217030 | |

| 217050 | 217050 | C6 deficiency |
|--|--------|--|
| 217070 | 217050 | |
| Conotruncal cardiac anomalies | 217070 | |
| 217300 | 217095 | |
| 21800 | | |
| 218000 Andermann syndrome 218030 Apparent mineralocorticoid excess, hypertension due to 221770 Polycystic lipomembranous osteodysplasia with sclerosing leukencephalopathy 221820 Gliosis, familial progressive subcortical 222100 Diabetes mellitus, insulin-dependent-1 222600 Achondrogenesis II, 256050 222600 Achondrogenesis II, 600972 222600 Diastrophic dysplasia 222700 Lysinuric protein intolerance 222800 Hemolytic anemia due to bisphosphoglycerate mutase deficiency 222900 Sucrose intolerance 2223900 Dysautonomia, familial 224100 Congenital dyserythropoietic anemia II 225500 Ellis-van Creveld syndrome 227220 [Eye color, brown] 227646 Fanconi anemia, type D 229300 Friedreich ataxia with retained reflexes 229800 [Fructosuria] 230000 Frucosidosis 230350 Galactose epimerase deficiency 230400 Galactosemia 230800 Gaucher disease with cardiovascular calcification 231550 Achalasia-addisonianism-alacrimia syndrome 231680 Glutaricaciduria, type II or pccB type 232200 Glycogen storage disease IIIa 232000 Glycogen storage disease IIIa 232000 Glycogen storage disease IIIa 232000 Glycogen storage disease IIIa 233710 Chronic granulomatous disease due to deficiency of NCF-2 234000 Factor XII deficiency 235200 Hemochromatosis [Histidinemia] 236100 Holoprosencephaly-1 | 217800 | |
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| 221820 Gliosis, familial progressive subcortical 222100 Diabetes mellitus, insulin-dependent-1 222600 Atelosteogenesis II, 256050 222600 Achondrogenesis II, 256050 222600 Diastrophic dysplasia 222700 Lysinuric protein intolerance 222800 Hemolytic anemia due to bisphosphoglycerate mutase deficiency 222900 Sucrose intolerance 223900 Dysautonomia, familial 224100 Congenital dyserythropoietic anemia II 225500 Ellis-van Creveld syndrome 227220 [Eye color, brown] 227646 Fanconi anemia, type D 229300 Friedreich ataxia 229300 Friedreich ataxia with retained reflexes 229800 Fructose intolerance 229800 [Fructosuria] 230000 Fucosidosis 230350 Galactose epimerase deficiency 230400 Galactose emimerase deficiency 230400 Gaucher disease 230800 Gaucher disease 230800 Gaucher disease 230800 Gaucher disease with cardiovascular calcification 231550 Achalasia-addisonianism-alacrimia syndrome 231680 Glutaricaciduria, type II or pccB type 232200 Glycogen storage disease IIIa 232400 Glycogen storage disease IIIb 233100 [Renal glucosuria] 233710 Chronic granulomatous disease due to deficiency of NCF-2 234000 Factor XII deficiency 235200 Hemochromatosis 235800 [Histidinemia] 236100 Holoprosencephaly-1 | | leukencenhalonathy |
| Diabetes mellitus, insulin-dependent-1 | 221820 | |
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| 222600 Achondrogenesis Ib, 600972 222600 Diastrophic dysplasia 222700 Lysinuric protein intolerance 222800 Hemolytic anemia due to bisphosphoglycerate mutase deficiency 222900 Sucrose intolerance 223900 Dysautonomia, familial 224100 Congenital dyserythropoietic anemia II 225500 Ellis-van Creveld syndrome 227220 [Eye color, brown] 227646 Fanconi anemia, type D 229300 Friedreich ataxia 229300 Friedreich ataxia with retained reflexes 229600 Fructose intolerance 229800 [Fructosuria] 230000 Fucosidosis 230350 Galactose epimerase deficiency 230400 Galactosemia 230800 Gaucher disease 230800 Gaucher disease 230800 Gaucher disease with cardiovascular calcification 231550 Achalasia-addisonianism-alacrimia syndrome 231680 Glutaricaciduria, type II or pccB type 232200 Glycogen storage disease IIIa 232400 Glycogen storage disease IIIa 232400 Glycogen storage disease IIIa 233100 [Renal glucosuria] 233710 Chronic granulomatous disease due to deficiency of NCF-2 234000 Factor XII deficiency 235200 Hemochromatosis 235800 [Histidinemia] 236100 Holoprosencephaly-1 | 222600 | |
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| Eructosuria Fucosidosis | 229600 | |
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| 231680 Glutaricaciduria, type IIA 232050 Propionicacidemia, type II or pccB type 232200 Glycogen storage disease I 232400 Glycogen storage disease IIIa 232400 Glycogen storage disease IIIb 233100 [Renal glucosuria] 233710 Chronic granulomatous disease due to deficiency of NCF-2 234000 Factor XII deficiency 235200 Hemochromatosis 235800 [Histidinemia] 236100 Holoprosencephaly-1 | 231550 | |
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| 233100 [Renal glucosuria] 233710 Chronic granulomatous disease due to deficiency of NCF-2 234000 Factor XII deficiency 235200 Hemochromatosis 235800 [Histidinemia] 236100 Holoprosencephaly-1 | 232400 | |
| 233100 [Renal glucosuria] 233710 Chronic granulomatous disease due to deficiency of NCF-2 234000 Factor XII deficiency 235200 Hemochromatosis 235800 [Histidinemia] 236100 Holoprosencephaly-1 | 232400 | Glycogen storage disease IIIb |
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| 235200 Hemochromatosis 235800 [Histidinemia] 236100 Holoprosencephaly-1 | 234000 | Factor XII deficiency |
| 235800 [Histidinemia] 236100 Holoprosencephaly-1 | 235200 | |
| | 235800 | [Histidinemia] |
| | 236100 | Holoprosencephaly-1 |
| | 236200 | Homocystinuria, B6-responsive and nonresponsive types |
| 236730 Urofacial syndrome | 236730 | Urofacial syndrome |

| 238310 | Hyperglycinemia, nonketotic, type II |
|----------|--|
| 239100 | Van Buchem disease |
| 240300 | Autoimmune polyglandular disease, type I |
| 243500 | Isovalericacidemia |
| 245050 | Ketoacidosis due to SCOT deficiency |
| 245200 | Krabbe disease |
| 245900 | Norum disease |
| 245900 | Fish-eye disease |
| 246450 | HMG-CoA lyase deficiency |
| 246530 | Leukotriene C4 synthase deficiency |
| 246900 | Lipoamide dehydrogenase deficiency |
| 247200 | Miller-Dieker lissencephaly syndrome |
| 248510 | Mannosidosis, beta- |
| 248600 | Maple syrup urine disease, type Ia |
| 248611 | Maple syrup urine disease, type Ib |
| 249000 | Meckel syndrome |
| 250100 | Metachromatic leukodystrophy |
| 250250 | Cartilage-hair hypoplasia |
| 250800 | Methemoglobinemia, type I |
| 250800 | Methemoglobinemia, type II |
| 251000 | Methylmalonicaciduria, mutase deficiency type |
| 251600 | Microphthalmia, autosomal recessive |
| 252920 . | Sanfilippo syndrome, type B |
| 252940 | Sanfilippo syndrome, type D |
| 253200 | Maroteaux-Lamy syndrome, several forms |
| 253250 | Mulibrey nanism |
| 253700 | Muscular dystrophy, limb-girdle, type 2C |
| 253800 | Walker-Warburg syndrome, 236670 |
| 253800 | Fukuyama type congenital muscular dystrophy |
| 254210 | Myasthenia gravis, familial infantile |
| 255800 | Schwartz-Jampel syndrome |
| 256100 | Nephronophthisis, juvenile |
| 256540 | Galactosialidosis |
| 256550 | Sialidosis, type I |
| 256550 | Sialidosis, type II |
| 257200 | Niemann-Pick disease, type A |
| 257200 | Niemann-Pick disease, type B |
| 258501 | 3-methylglutaconicaciduria, type III |
| 258870 | Gyrate atrophy of choroid and retina with ornithinemia, B6 |
| | responsive or unresponsive |
| 261510 | Pseudo-Zellweger syndrome |
| 261600 | Phenylketonuria |
| 261600 | [Hyperphenylalaninemia, mild] |
| 261640 | Phenylketonuria due to PTS deficiency |
| 263200 | Polycystic kidney disease, autosomal recessive |
| 263700 | Porphyria, congenital erythropoietic |

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| 264300 | Pseudohermaphroditism, male, with gynecomastia |
| 264700 | Pseudo-vitamin D dependency rickets 1 |
| 266200 | Anemia, hemolytic, due to PK deficiency |
| 266600 | Inflammatory bowel disease-1 |
| 267750 | Knobloch syndrome |
| 268800 | Sandhoff disease, infantile, juvenile, and adult forms |
| 268800 | Spinal muscular atrophy, HEXB-related |
| 269920 | Salla disease |
| 270100 | Situs inversus viscerum |
| 271245 | Spinocerebellar ataxia-8, infantile, with sensory neuropathy |
| 271900 | Canavan disease |
| 272750 | GM2-gangliosidosis, AB variant |
| 272800 | Tay-Sachs disease |
| 272800 | [Hex A pseudodeficiency] |
| 272800 | GM2-gangliosidosis, juvenile, adult |
| 273300 | Male germ cell tumor |
| 274600 | Pendred syndrome |
| 274600 | Deafness, autosomal recessive 4 |
| 276600 | Tyrosinemia, type II |
| 276700 | Tyrosinemia, type I |
| 276900 | Usher syndrome, type 1A |
| 276902 | Usher syndrome, type 3 |
| 278000 | Wolman disease |
| 278000 | Cholesteryl ester storage disease |
| 278300 | Xanthinuria, type I |
| 278760 | Xeroderma pigmentosum, group F |
| 300031 | Mental retardation, X-linked, FRAXF type |
| 300044 | Wernicke-Korsakoff syndrome, susceptibility to |
| 300048 | Intestinal pseudoobstruction, neuronal, X-linked |
| 300049 | Nodular heterotopia, bilateral periventricular |
| 300049 | BPNH/MR syndrome |
| 300055 | Mental retardation with psychosis, pyramidal signs, and |
| | macroorchidism |
| 300100 | Adrenoleukodystrophy |
| 300100 | Adrenomyeloneuropathy |
| 300104 | Mental retardation, X-linked nonspecific, 309541 |
| 300126 | Dyskeratosis congenita-1, 305000 |
| 301201 | Amelogenesis imperfecta-3, hypoplastic type |
| 301220 | Partington syndrome II |
| 301590 | Anophthalmos-1 |
| 302060 | Noncompaction of left ventricular myocardium, isolated |
| 302060 | Barth syndrome |
| 302060 | Cardiomyopathy, X-linked dilated, 300069 |
| 302060 | Endocardial fibroelastosis-2 |
| 302350 | Nance-Horan syndrome |
| 302960 | Chondrodysplasia punctata, X-linked dominant |
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| 303700 | Colorblindness, blue monochromatic |
|--------|--|
| 303800 | Colorblindness, deutan |
| 303900 | Colorblindness, protan |
| 304040 | Charcot-Marie-Tooth neuropathy, X-linked-1, dominant, 302800 |
| 304800 | Diabetes insipidus, nephrogenic |
| 305100 | Anhidrotic ectodermal dysplasia |
| 305450 | FG syndrome |
| 305900 | Favism |
| 305900 | G6PD deficiency |
| 305900 | Hemolytic anemia due to G6PD deficiency |
| 306400 | Chronic granulomatous disease, X-linked |
| 306700 | Hemophilia A |
| 306995 | [Homosexuality, male] |
| 308310 | Incontinentia pigmenti, familial |
| 308840 | Spastic paraplegia, 312900 |
| 308840 | Hydrocephalus due to aqueductal stenosis, 307000 |
| 308840 | MASA syndrome, 303350 |
| 309200 | Manic-depressive illness, X-linked |
| 309470 | Mental retardation, X-linked, syndromic-3, with spastic diplegia |
| 309548 | Mental retardation, X-linked, FRAXE type |
| 309585 | Mental retardation, X-linked, syndromic-6, with gynecomastia |
| | and obesity |
| 309600 | Allan-Herndon syndrome |
| 309605 | Mental retardation, X-linked, syndromic-4, with congenital |
| | contractures and low fingertip arches |
| 309620 | Mental retardation-skeletal dysplasia |
| 309900 | Mucopolysaccharidosis II |
| 310300 | Emery-Dreifuss muscular dystrophy |
| 310400 | Myotubular myopathy, X-linked |
| 310460 | Myopia-1 |
| 310460 | Bornholm eye disease |
| 311250 | Ornithine transcarbamylase deficiency |
| 311300 | Otopalatodigital syndrome, type I |
| 311360 | Ovarian failure, premature |
| 311510 | Waisman parkinsonism-mental retardation syndrome |
| 312040 | N syndrome, 310465 |
| 312610 | Retinitis pigmentosa-3 |
| 312760 | Turner syndrome |
| 314250 | Dystonia-3, torsion, with parkinsonism, Filipino type |
| 314300 | Goeminne TKCR syndrome |
| 314400 | Cardiac valvular dysplasia-1 |
| 314580 | Wieacker-Wolff syndrome |
| 314850 | McLeod phenotype |
| 400003 | Sertoli-cell-only syndrome |
| 415000 | Sertoli-cell-only syndrome |
| 600020 | Prostate cancer, 176807 |
| | |

| 600040 | |
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| 600040 | Colorectal cancer |
| 600049 | Myelodysplasia syndrome-1 |
| 600059 | Retinitis pigmentosa-13 |
| 600065 | Leukocyte adhesion deficiency, 116920 |
| 600095 | Split hand/foot malformation, type 3 |
| 600101 | Deafness, autosomal dominant 2 |
| 600105 | Retinitis pigmentosa-12, autosomal recessive |
| 600119 | Muscular dystrophy, Duchenne-like, type 2 |
| 600119 | Adhalinopathy, primary |
| 600138 | Retinitis pigmentosa-11 |
| 600163 | Long QT syndrome-3 |
| 600173 | SCID, autosomal recessive, T-negative/B-positive type |
| 600175 | Spinal muscular atrophy, congenital nonprogressive, of lower limbs |
| 600179 | Leber congenital amaurosis, type I, 204000 |
| 600184 | Carnitine acetyltransferase deficiency |
| 600194 | Ichthyosis bullosa of Siemens, 146800 |
| 600202 | Dyslexia, specific, 2 |
| 600211 | Cleidocranial dysplasia, 119600 |
| 600223 | Spinocerebellar ataxia-4 |
| 600231 | Palmoplantar keratoderma, Bothnia type |
| 600243 | Temperature-sensitive apoptosis |
| 600261 | Ehlers-Danlos-like syndrome |
| 600276 | Cerebral arteriopathy with subcortical infarcts and |
| | leukoencephalopathy, 125310 |
| 600281 | Non-insulin-dependent diabetes mellitus, 125853 |
| 600281 | MODY, type 1, 125850 |
| 600309 | Atrioventricular canal defect-1 |
| 600310 | Pseudoachondroplasia, 177170 |
| 600310 | Epiphyseal dysplasia, multiple 1, 132400 |
| 600320 | Insulin-dependent diabetes mellitus-5 |
| 600334 | Tibial muscular dystrophy |
| 600354 | Spinal muscular atrophy-1, 253300 |
| 600354 | Spinal muscular atrophy-2, 253550 |
| 600354 | Spinal muscular atrophy-3, 253400 |
| 600374 | Bardet-Biedl syndrome 4 |
| 600429 | [Ii blood group, 110800] |
| 600512 | Epilepsy, partial |
| 600536 | Myopathy, congenital |
| 600542 | Chondrosarcoma, extraskeletal myxoid |
| 600584 | Atrial septal defect with atrioventricular conduction defects, |
| 600593 | Craniosynostosis, Adelaide type |
| 600650 | Myopathy due to CPT II deficiency, 255110 |
| 600650 | CPT deficiency, hepatic, type II, 600649 |
| 600652 | Deafness, autosomal dominant 4 |
| 10001 | Doumoss, autosomai uommant 4 |

| 600698 | Salivary adenoma |
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| 600698 | Uterine leiomyoma |
| 600698 | Lipoma |
| 600698 | Lipomatosis, mutiple, 151900 |
| 600701 | Lipoma |
| 600722 | Ceroid lipofuscinosis, neuronal, variant juvenile type, with |
| | granular osmiophilic deposits |
| 600722 | Ceroid lipofuscinosis, neuronal-1, infantile, 256730 |
| 600757 | Orofacial cleft-3 |
| 600759 | Alzheimer disease-4 |
| 600760 | Pseudohypoaldosteronism, type I, 264350 |
| 600760 | Liddle syndrome, 177200 |
| 600761 | Pseudohypoaldosteronism, type I, 264350 |
| 600761 | Liddle syndrome, 177200 |
| 600807 | Bronchial asthma |
| 600808 | Enuresis, nocturnal, 2 |
| 600837 | Hirschsprung disease, 142623 |
| 600839 | Bartter syndrome, 241200 |
| 600850 | Schizophrenia disorder-4 |
| 600856 | Beckwith-Wiedemann syndrome, 130650 |
| 600881 | Cataract, congenital, zonular, with sutural opacities |
| 600882 | Charcot-Marie-Tooth neuropathy-2B |
| 600883 | Diabetes mellitus, insulin-dependent, 8 |
| 600884 | Cardiomyopathy, familial dilated 1B |
| 600887 | Endometrial carcinoma |
| 600897 | Cataract, zonular pulverulent-1, 116200 |
| 600899 | Severe combined immunodeficiency, type I, 202500 |
| 600918 | Cystinuria, type III |
| 600919 | Long QT syndrome-4 with sinus bradycardia |
| 600923 | Porphyria variegata, 176200 |
| 600946 | Short stature, autosomal dominant, with normal serum growth |
| | hormone binding protein |
| 600946 | Short stature, idiopathic |
| 600946 | Laron dwarfism, 262500 |
| 600956 | Persistent Mullerian duct syndrome, type II 261550 |
| 600957 | Persistent Mullerian duct syndrome, type I 261550 |
| 600968 | Gitelman syndrome, 263800 |
| 600971 | Deafness, autosomal recessive 6 |
| 600974 | Deafness, autosomal recessive 7 |
| 600977 | Cone dystrophy, progressive |
| 600994 | Deafness, autosomal dominant 5 |
| 600995 | Nephrotic syndrome, idiopathic, steroid-resistant |
| 600998 | Bleeding diathesis due to GNAQ deficiency |
| 601002 | 5-oxoprolinuria, 266130 |
| 601002 | Hemolytic anemia due to glutathione synthetase deficiency, 231900 |

| 601071 | Deafness, autosomal recessive 9 |
|---------|---|
| 601072 | Deafness, autosomal recessive 8 |
| 601105 | Pycnodysostosis, 265800 |
| 601107 | Dubin-Johnson syndrome, 237500 |
| 601130 | Tolbutamide poor metabolizer |
| 601145 | Epilepsy, progressive myoclonic 1, 254800 |
| 601146 | Brachydactyly, type C, 113100 |
| 601146 | Acromesomelic dysplasia, Hunter-Thompson type, 201250 |
| 601146 | Chondrodysplasia, Grebe type, 200700 |
| 601154 | Cardiomyopathy, dilated, 1E |
| 601199 | Neonatal hyperparathyroidism, 239200 |
| 601199 | Hypocalcemia, autosomal dominant, 601198 |
| 601199 | Hypocalciuric hypercalcemia, type I, 145980 |
| 601202 | Cataract, anterior polar-2 |
| 601208 | Insulin-dependent diabetes mellitus-11 |
| 601226 | Progressive external ophthalmoplegia, type 2 |
| 601238 | Cerebellar ataxia, Cayman type |
| 601267 | HIV infection, susceptibility/resistence to |
| 601284 | Hereditary hemorrhagic telangiectasia-2, 600376 |
| 601313 | Polycystic kidney disease, adult type I, 173900 |
| 601316 | Deafness, autosomal dominant 10 |
| 601363 | Wilms tumor, type 4 |
| 601373 | HIV infection, susceptibility/resistance to |
| 601402 | Leukemia, myeloid, acute |
| 601406 | B-cell non-Hodgkin lymphoma, high-grade |
| 601410 | Diabetes mellitus, transient neonatal |
| 601411 | Muscular dystrophy, limb-girdle, type 2F, 601287 |
| 601412 | Deafness, autosomal dominant 7 |
| 601414 | Retinitis pigmentosa-18 |
| 601471· | Moebius syndrome-2 |
| 601493 | Cardiomyopathy, dilated 1C |
| 601494 | Cardiomyopathy, familial, dilated-2 |
| 601518 | Prostate cancer, hereditary, 1, 176807 |
| 601542 | Rieger syndrome, type 1, 180500 |
| 601545 | Lissencephaly-1 |
| 601596 | Charcot-Marie-Tooth neuropathy, demyelinating |
| 601604 | Mycobacterial and salmonella infections, susceptibility to |
| 601607 | Rhabdoid tumors |
| 601620 | Holt-Oram syndrome, 142900 |
| 601621 | Ulnar-mammary syndrome, 181450 |
| 601649 | Blepharophimosis, epicanthus inversus, and ptosis, type 2 |
| 601652 | Glaucoma 1A, primary open angle, juvenile-onset, 137750 |
| 601669 | Hirschsprung disease, one form |
| 601680 | Distal arthrogryposis, type 2B |
| 601682 | C1 1.C : |
| 601690 | Glaucoma 1C, primary open angle Platelet-activating factor acetylhydrolase deficiency |

| (01/01 | D-4:-4:- 10 (01710 |
|--------|---|
| 601691 | Retinitis pigmentosa-19, 601718 |
| 601691 | Stargardt disease-1, 248200 |
| 601691 | Cone-rod dystrophy 3 |
| 601691 | Fundus flavimaculatus with macular dystrophy, 248200 |
| 601718 | Retinitis pigmentosa-19 |
| 601757 | Rhizomelic chondrodysplasia punctata, type 1, 215100 |
| 601769 | Osteoporosis, involutional |
| 601769 | Rickets, vitamin D-resistant, 277440 |
| 601771 | Glaucoma 3A, primary infantile, 231300 |
| 601777 | Cone dystrophy, progressive |
| 601780 | Ceroid-lipofuscinosis, neuronal-6, variant late infantile |
| 601800 | [Hair color, brown] |
| 601841 | Protein C inhibitor deficiency |
| 601843 | Hypothyroidism, congenital, 274400 |
| 601844 | Pseudohypoaldosteronism type II |
| 601846 | Muscular dystrophy with rimmed vacuoles |
| 601863 | Bare lymphocyte syndrome, complementation group C |
| 601868 | Deafness, autosomal dominant 13 |
| 601885 | Cataract, zonular pulverulent-2 |
| 601916 | Pancreatic cancer |
| 601928 | Monilethrix, 158000 |
| 601954 | Muscular dystrophy, limb-girdle, type 2G |
| 601969 | Glioblastoma multiforme, 137800 |
| 601969 | Medulloblastoma, 155255 |
| 601975 | Ectodermal dysplasia/skin fragility syndrome |
| 602014 | Hypomagnesemia with secondary hypocalcemia |
| 602066 | Convulsions, infantile and paroxysmal choreoathetosis |
| 602067 | Cardiomyopathy, dilated, 1F |
| 602081 | Speech-language disorder-1 |
| 602082 | Corneal dystrophy, Thiel-Behnke type |
| 602084 | Endometrial carcinoma |
| 602088 | Nephronophthisis, infantile |
| 602089 | Hemangioma, capillary, hereditary |
| 602092 | Deafness, autosomal recessive 18 |
| 602094 | Lipodystrophy, familial partial |
| 602116 | Glioma |
| 602134 | Tremor, familial essential, 2 |
| 602136 | Refsum disease, infantile, 266510 |
| 602136 | Zellweger syndrome-1, 214100 |
| 602136 | Adrenoleukodystrophy, neonatal, 202370 |
| 602153 | Monilethrix, 158000 |
| 602216 | Peutz-Jeghers syndrome, 175200 |
| 602221 | Stem-cell leukemia/lymphoma syndrome |
| 602225 | Cone-rod retinal dystrophy-2, 120970 |
| 602225 | Leber congenital amaurosis, type III |
| 602279 | Oculopharyngeal muscular dystorphy, 164300 |
| | 1 |

| 602279 | Oculopharyngeal muscular dystrophy, autosomal recessive, 257950 |
|--------|---|
| 602280 | Retinitis pigmentosa-14, 600132 |
| 602363 | Ellis-van Creveld-like syndrome |
| 602403 | Alzheimer disease, susceptibility to |
| 602421 | Sweat chloride elevation without CF |
| 602421 | Congenital bilateral absence of vas deferens, 277180 |
| 602421 | Cystic fibrosis, 219700 |
| 602447 | Coronary artery disease, susceptibility to |
| 602475 | Ossification of posterior longitudinal ligament of spine |
| 602476 | Febrile convulsions, familial, 1 |
| 602477 | Febrile convulsions, familial, 2 |
| 602491 | Hyperlipidemia, familial combined, 1 |
| 602544 | Parkinson disease, juvenile, type 2, 600116 |
| 602568 | Homocystinuria-megaloblastic anemia, cbl E type, 236270 |
| 602574 | Deafness, autosomal dominant 12, 601842 |
| 602574 | Deafness, autosomal dominant 8, 601543 |
| 602575 | Nail-patella syndrome with open-angle glaucoma, 137750 |
| 602575 | Nail-patella syndrome, 161200 |
| 602629 | Dystonia-6, torsion |
| 602631 | Rhabdomyosarcoma, 268210 |
| 602631 | Breast Cancer |
| 602669 | Anterior segment mesenchymal dysgenesis and cataract, 107250 |
| 602669 | Cataract, congenital |
| 602685 | Mental retardation, severe, with spasticity and tapetoretinal |
| | degeneration |
| 602716 | Nephrosis-1, congenital, Finnish type, 256300 |
| 602771 | Muscular dystrophy, congenital, with early spine rigidity |
| 602772 | Retinitis pitmentosa-24 |

Polynucleotide and Polypeptide Variants

The present invention is also directed to variants of the immune/hematopoietic associated polynucleotide sequence disclosed in SEQ ID NO:X or the complementary strand thereto, nucleotide sequences encoding the polypeptide of SEQ ID NO:Y, the nucleotide sequence of SEQ ID NO:X encoding the polypeptide sequence as defined in column 6 of Table 1A, nucleotide sequences encoding the polypeptide as defined in column 6 of Table 1A, the nucleotide sequence as defined in columns 8 and 9 of Table 2, nucleotide sequences encoding the polypeptide encoded by the nucleotide sequence as defined in columns 8 and 9 of Table 2, the nucleotide sequence as defined in column 6 of Table 1B, nucleotide sequences encoding the polypeptide encoded by the nucleotide sequence as defined in column 6 of Table 1B, the cDNA sequence contained in Clone ID NO:Z, and/or nucleotide sequences encoding a polypeptide encoded by the cDNA sequence contained in Clone ID NO:Z.

- [092] The present invention also encompasses variants of the polypeptide sequence disclosed in SEQ ID NO:Y, a polypeptide sequence as defined in column 6 of Table 1A, a polypeptide sequence encoded by the polynucleotide sequence in SEQ ID NO:X, a polypeptide sequence encoded by the nucleotide sequence as defined in columns 8 and 9 of Table 2, a polypeptide sequence encoded by the nucleotide sequence as defined in column 6 of Table 1B, a polypeptide sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X, and/or a polypeptide sequence encoded by the cDNA sequence contained in Clone ID NO:Z.
- [093] "Variant" refers to a polynucleotide or polypeptide differing from the polynucleotide or polypeptide of the present invention, but retaining essential properties thereof. Generally, variants are overall closely similar, and, in many regions, identical to the polynucleotide or polypeptide of the present invention.
- Thus, one aspect of the invention provides an isolated nucleic acid molecule comprising, or alternatively consisting of, a polynucleotide having a nucleotide sequence selected from the group consisting of: (a) a nucleotide sequence described in SEQ ID NO:X or contained in the cDNA sequence of Clone ID NO:Z; (b) a nucleotide sequence in SEQ ID NO:X or the cDNA in Clone ID NO:Z which encodes a mature immune/hematopoietic associated polypeptide; (c) a nucleotide sequence in SEQ ID NO:X or the cDNA sequence of Clone ID NO:Z, which encodes a biologically active

fragment of an immune/hematopoietic associated polypeptide; (d) a nucleotide sequence in SEQ ID NO:X or the cDNA sequence of Clone ID NO:Z, which encodes an antigenic fragment of an immune/hematopoietic associated polypeptide; (e) a nucleotide sequence encoding an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; (f) a nucleotide sequence encoding a mature immune/hematopoietic associated polypeptide of the amino acid sequence of SEO ID NO:Y or the amino acid sequence encoded by the cDNA in Clone ID NO:Z; (g) a nucleotide sequence encoding a biologically active fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; (h) a nucleotide sequence encoding an antigenic fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; and (i) a nucleotide sequence complementary to any of the nucleotide sequences in (a), (b), (c), (d), (e), (f), (g), or (h), above.

[095] The present invention is also directed to nucleic acid molecules which comprise, or alternatively consist of, a nucleotide sequence which is at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100%, identical to, for example, any of the nucleotide sequences in (a), (b), (c), (d), (e), (f), (g), (h), or (i) above, the nucleotide coding sequence in SEQ ID NO:X or the complementary strand thereto, the nucleotide coding sequence of the cDNA contained in Clone ID NO:Z or the complementary strand thereto, a nucleotide sequence encoding the polypeptide of SEQ ID NO:Y, a nucleotide sequence encoding a polypeptide sequence encoded by the nucleotide sequence in SEQ ID NO:X, a polypeptide sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X, a nucleotide sequence encoding the polypeptide encoded by the cDNA contained in Clone ID NO:Z, the nucleotide coding sequence in SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto, a nucleotide sequence encoding the polypeptide encoded by the nucleotide sequence in SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto, the nucleotide coding sequence in SEQ ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto, a nucleotide sequence encoding the polypeptide encoded by the nucleotide sequence in SEO

ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto, the nucleotide sequence in SEQ ID NO:X encoding the polypeptide sequence as defined in column 6 of Table 1A or the complementary strand thereto, nucleotide sequences encoding a polypeptide as defined in column 6 of Table 1A or the complementary strand thereto, and/or polynucleotide fragments of any of these nucleic acid molecules (e.g., those fragments described herein). Polynucleotides which hybridize to the complement of these nucleic acid molecules under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are polypeptides encoded by these polynucleotides and nucleic acids.

[096] In a preferred embodiment, the invention encompasses nucleic acid molecules which comprise, or alternatively, consist of a polynucleotide which hybridizes under stringent hybridization conditions, or alternatively, under lower stringency conditions, to a polynucleotide in (a), (b), (c), (d), (e), (f), (g), (h), or (i) above, as are polypeptides encoded by these polynucleotides. In another preferred embodiment, polynucleotides which hybridize to the complement of these nucleic acid molecules under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

In another embodiment, the invention provides a purified protein comprising, or alternatively consisting of, a polypeptide having an amino acid sequence selected from the group consisting of: (a) the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; (b) the amino acid sequence of a mature immune/hematopoietic associated polypeptide having the amino acid sequence of SEQ ID NO:Y or the amino acid sequence encoded by the cDNA in Clone ID NO:Z; (c) the amino acid sequence of a biologically active fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; and (d) the amino acid sequence of an antigenic fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z.

[098] The present invention is also directed to proteins which comprise, or alternatively consist of, an amino acid sequence which is at least 80%, 85%, 90%, 95%,

96%, 97%, 98%, 99% or 100%, identical to, for example, any of the amino acid sequences in (a), (b), (c), or (d), above, the amino acid sequence shown in SEQ ID NO:Y, the amino acid sequence encoded by the cDNA contained in Clone ID NO:Z, the amino acid sequence of the polypeptide encoded by the nucleotide sequence in SEQ ID NO:X as defined in columns 8 and 9 of Table 2, the amino acid sequence of the polypeptide encoded by the nucleotide sequence in SEQ ID NO:B as defined in column 6 of Table 1B, the amino acid sequence as defined in column 6 of Table 1A, an amino acid sequence encoded by the nucleotide sequence in SEQ ID NO:X, and an amino acid sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X. Fragments of these polypeptides are also provided (e.g., those fragments described herein). Further proteins encoded by polynucleotides which hybridize to the complement of the nucleic acid molecules encoding these amino acid sequences under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are the polynucleotides encoding these proteins.

"identical" to a reference nucleotide sequence of the present invention, it is intended that the nucleotide sequence of the nucleic acid is identical to the reference sequence except that the nucleotide sequence may include up to five point mutations per each 100 nucleotides of the reference nucleotide sequence encoding the polypeptide. In other words, to obtain a nucleic acid having a nucleotide sequence at least 95% identical to a reference nucleotide sequence, up to 5% of the nucleotides in the reference sequence may be deleted or substituted with another nucleotide, or a number of nucleotides up to 5% of the total nucleotides in the reference sequence may be inserted into the reference sequence. The query sequence may be an entire sequence referred to in Table 1A or 2 as the ORF (open reading frame), or any fragment specified, as described herein.

[0100] As a practical matter, whether any particular nucleic acid molecule or polypeptide is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a nucleotide sequence of the present invention can be determined conventionally using known computer programs. A preferred method for determining the best overall match between a query sequence (a sequence of the present invention) and a subject sequence, also referred to as a global sequence alignment, can be determined using the FASTDB computer program based on the algorithm of Brutlag et al. (Comp. App. Biosci. 6:237-245

(1990)). In a sequence alignment the query and subject sequences are both DNA sequences. An RNA sequence can be compared by converting U's to T's. The result of said global sequence alignment is expressed as percent identity. Preferred parameters used in a FASTDB alignment of DNA sequences to calculate percent identity are: Matrix=Unitary, k-tuple=4, Mismatch Penalty=1, Joining Penalty=30, Randomization Group Length=0, Cutoff Score=1, Gap Penalty=5, Gap Size Penalty 0.05, Window Size=500 or the length of the subject nucleotide sequence, whichever is shorter.

If the subject sequence is shorter than the query sequence because of 5' or 3' [0101]deletions, not because of internal deletions, a manual correction must be made to the results. This is because the FASTDB program does not account for 5' and 3' truncations of the subject sequence when calculating percent identity. For subject sequences truncated at the 5' or 3' ends, relative to the query sequence, the percent identity is corrected by calculating the number of bases of the query sequence that are 5' and 3' of the subject sequence, which are not matched/aligned, as a percent of the total bases of the query sequence. Whether a nucleotide is matched/aligned is determined by results of the FASTDB sequence alignment. This percentage is then subtracted from the percent identity, calculated by the above FASTDB program using the specified parameters, to arrive at a final percent identity score. This corrected score is what is used for the purposes of the present invention. Only bases outside the 5' and 3' bases of the subject sequence, as displayed by the FASTDB alignment, which are not matched/aligned with the query sequence, are calculated for the purposes of manually adjusting the percent identity score.

sequence to determine percent identity. The deletions occur at the 5' end of the subject sequence and therefore, the FASTDB alignment does not show a matched/alignment of the first 10 bases at 5' end. The 10 unpaired bases represent 10% of the sequence (number of bases at the 5' and 3' ends not matched/total number of bases in the query sequence) so 10% is subtracted from the percent identity score calculated by the FASTDB program. If the remaining 90 bases were perfectly matched the final percent identity would be 90%. In another example, a 90 base subject sequence is compared with a 100 base query sequence. This time the deletions are internal deletions so that there are no bases on the 5' or 3' of the subject sequence which are not matched/aligned with the query. In this case

the percent identity calculated by FASTDB is not manually corrected. Once again, only bases 5' and 3' of the subject sequence which are not matched/aligned with the query sequence are manually corrected for. No other manual corrections are to be made for the purposes of the present invention.

"identical" to a query amino acid sequence of the present invention, it is intended that the amino acid sequence of the subject polypeptide is identical to the query sequence except that the subject polypeptide sequence may include up to five amino acid alterations per each 100 amino acids of the query amino acid sequence. In other words, to obtain a polypeptide having an amino acid sequence at least 95% identical to a query amino acid sequence, up to 5% of the amino acid residues in the subject sequence may be inserted, deleted, (indels) or substituted with another amino acid. These alterations of the reference sequence may occur at the amino or carboxy terminal positions of the reference amino acid sequence or anywhere between those terminal positions, interspersed either individually among residues in the reference sequence or in one or more contiguous groups within the reference sequence.

[0104] As a practical matter, whether any particular polypeptide is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance, the amino acid sequence of a polypeptide referred to in Table 1A (e.g., an amino acid sequence identified in columns 5 or 6) or Table 2 (e.g., the amino acid sequence of the polypeptide encoded by the polynucleotide sequence defined in columns 8 and 9 of Table 2) or a fragment thereof, the amino acid sequence of the polypeptide encoded by the polynucleotide sequence in SEQ ID NO:B as defined in column 6 of Table 1B or a fragment thereof, the amino acid sequence of the polypeptide encoded by the nucleotide sequence in SEQ ID NO:X or a fragment thereof, or an amino acid sequence of the polypeptide encoded by cDNA contained in Clone ID NO:Z, or a fragment thereof, can be determined conventionally using known computer programs. A preferred method for determining the best overall match between a query sequence (a sequence of the present invention) and a subject sequence, also referred to as a global sequence alignment, can be determined using the FASTDB computer program based on the algorithm of Brutlag et al. (Comp. App. Biosci.6:237-245 (1990)). In a sequence alignment the query and subject sequences are either both nucleotide sequences or both amino acid sequences. The result of said global

sequence alignment is expressed as percent identity. Preferred parameters used in a FASTDB amino acid alignment are: Matrix=PAM 0, k-tuple=2, Mismatch Penalty=1, Joining Penalty=20, Randomization Group Length=0, Cutoff Score=1, Window Size=sequence length, Gap Penalty=5, Gap Size Penalty=0.05, Window Size=500 or the length of the subject amino acid sequence, whichever is shorter.

If the subject sequence is shorter than the query sequence due to N- or C-[0105]terminal deletions, not because of internal deletions, a manual correction must be made to the results. This is because the FASTDB program does not account for N- and C-terminal truncations of the subject sequence when calculating global percent identity. For subject sequences truncated at the N- and C-termini, relative to the query sequence, the percent identity is corrected by calculating the number of residues of the query sequence that are N- and C-terminal of the subject sequence, which are not matched/aligned with a corresponding subject residue, as a percent of the total bases of the query sequence. Whether a residue is matched/aligned is determined by results of the FASTDB sequence alignment. This percentage is then subtracted from the percent identity, calculated by the above FASTDB program using the specified parameters, to arrive at a final percent identity score. This final percent identity score is what is used for the purposes of the present invention. Only residues to the N- and C-termini of the subject sequence, which are not matched/aligned with the query sequence, are considered for the purposes of manually adjusting the percent identity score. That is, only query residue positions outside the farthest N- and C- terminal residues of the subject sequence.

residue query sequence to determine percent identity. The deletion occurs at the N-terminus of the subject sequence and therefore, the FASTDB alignment does not show a matching/alignment of the first 10 residues at the N-terminus. The 10 unpaired residues represent 10% of the sequence (number of residues at the N- and C- termini not matched/total number of residues in the query sequence) so 10% is subtracted from the percent identity score calculated by the FASTDB program. If the remaining 90 residues were perfectly matched the final percent identity would be 90%. In another example, a 90 residue subject sequence is compared with a 100 residue query sequence. This time the deletions are internal deletions so there are no residues at the N- or C-termini of the subject sequence which are not matched/aligned with the query. In this case the percent

identity calculated by FASTDB is not manually corrected. Once again, only residue positions outside the N- and C-terminal ends of the subject sequence, as displayed in the FASTDB alignment, which are not matched/aligned with the query sequence are manually corrected for. No other manual corrections are to be made for the purposes of the present invention.

- [0107] The polynucleotide variants of the invention may contain alterations in the coding regions, non-coding regions, or both. Especially preferred are polynucleotide variants containing alterations which produce silent substitutions, additions, or deletions, but do not alter the properties or activities of the encoded polypeptide. Nucleotide variants produced by silent substitutions due to the degeneracy of the genetic code are preferred. Moreover, polypeptide variants in which less than 50, less than 40, less than 30, less than 20, less than 10, or 5-50, 5-25, 5-10, 1-5, or 1-2 amino acids are substituted, deleted, or added in any combination are also preferred. Polynucleotide variants can be produced for a variety of reasons, e.g., to optimize codon expression for a particular host (change codons in the human mRNA to those preferred by a bacterial host such as *E. coli*).
- [0108] Naturally occurring variants are called "allelic variants," and refer to one of several alternate forms of a gene occupying a given locus on a chromosome of an organism. (Genes II, Lewin, B., ed., John Wiley & Sons, New York (1985).) These allelic variants can vary at either the polynucleotide and/or polypeptide level and are included in the present invention. Alternatively, non-naturally occurring variants may be produced by mutagenesis techniques or by direct synthesis.
- [0109] Using known methods of protein engineering and recombinant DNA technology, variants may be generated to improve or alter the characteristics of the polypeptides of the present invention. For instance, one or more amino acids can be deleted from the N-terminus or C-terminus of the polypeptides of the present invention without substantial loss of biological function. As an example, the authors of Ron et al., J. Biol. Chem. 268: 2984-2988 (1993), reported variant KGF proteins having heparin binding activity even after deleting 3, 8, or 27 amino-terminal amino acid residues. Similarly, Interferon gamma exhibited up to ten times higher activity after deleting 8-10 amino acid residues from the carboxy terminus of this protein. (Dobeli et al., J. Biotechnology 7:199-216 (1988).)

[0110] Moreover, ample evidence demonstrates that variants often retain a biological activity similar to that of the naturally occurring protein. For example, Gayle and coworkers (J. Biol. Chem. 268:22105-22111 (1993)) conducted extensive mutational analysis of human cytokine IL-1a. They used random mutagenesis to generate over 3,500 individual IL-1a mutants that averaged 2.5 amino acid changes per variant over the entire length of the molecule. Multiple mutations were examined at every possible amino acid position. The investigators found that "[m]ost of the molecule could be altered with little effect on either [binding or biological activity]." In fact, only 23 unique amino acid sequences, out of more than 3,500 nucleotide sequences examined, produced a protein that significantly differed in activity from wild-type.

- [0111] Furthermore, even if deleting one or more amino acids from the N-terminus or C-terminus of a polypeptide results in modification or loss of one or more biological functions, other biological activities may still be retained. For example, the ability of a deletion variant to induce and/or to bind antibodies which recognize the secreted form will likely be retained when less than the majority of the residues of the secreted form are removed from the N-terminus or C-terminus. Whether a particular polypeptide lacking N-or C-terminal residues of a protein retains such immunogenic activities can readily be determined by routine methods described herein and otherwise known in the art.
- [0112] Thus, the invention further includes polypeptide variants which show a functional activity (e.g., biological activity) of the polypeptides of the invention. Such variants include deletions, insertions, inversions, repeats, and substitutions selected according to general rules known in the art so as have little effect on activity.
- [0113] The present application is directed to nucleic acid molecules at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100% identical to the nucleic acid sequences disclosed herein, (e.g., encoding a polypeptide having the amino acid sequence of an N and/or C terminal deletion), irrespective of whether they encode a polypeptide having functional activity. This is because even where a particular nucleic acid molecule does not encode a polypeptide having functional activity, one of skill in the art would still know how to use the nucleic acid molecule, for instance, as a hybridization probe or a polymerase chain reaction (PCR) primer. Uses of the nucleic acid molecules of the present invention that do not encode a polypeptide having functional activity include, *interalia*, (1) isolating a gene or allelic or splice variants thereof in a cDNA library; (2) in situ

hybridization (e.g., "FISH") to metaphase chromosomal spreads to provide precise chromosomal location of the gene, as described in Verma et al., Human Chromosomes: A Manual of Basic Techniques, Pergamon Press, New York (1988); (3) Northern Blot analysis for detecting mRNA expression in specific tissues (e.g., normal immune/hematopoietic or diseased immune/hematopoietic tissues); and (4) in situ hybridization (e.g., histochemistry) for detecting mRNA expression in specific tissues (e.g., normal immune/hematopoietic or diseased immune/hematopoietic tissues).

[0114] Preferred, however, are nucleic acid molecules having sequences at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100% identical to the nucleic acid sequences disclosed herein, which do, in fact, encode a polypeptide having functional activity. By a polypeptide having "functional activity" is meant, a polypeptide capable of displaying one or more known functional activities associated with a full-length (complete) protein of the invention. Such functional activities include, but are not limited to, biological activity, antigenicity [ability to bind (or compete with a polypeptide of the invention for binding) to an anti-polypeptide of the invention antibody], immunogenicity (ability to generate antibody which binds to a specific polypeptide of the invention), ability to form multimers with polypeptides of the invention, and ability to bind to a receptor or ligand for a polypeptide of the invention.

[0115] The functional activity of the polypeptides, and fragments, variants and derivatives of the invention, can be assayed by various methods.

or compete with full-length polypeptide of the present invention for binding to an antipolypeptide of the invention antibody, various immunoassays known in the art can be
used, including but not limited to, competitive and non-competitive assay systems using
techniques such as radioimmunoassays, ELISA (enzyme linked immunosorbent assay),
"sandwich" immunoassays, immunoradiometric assays, gel diffusion precipitation
reactions, immunodiffusion assays, in situ immunoassays (using colloidal gold, enzyme or
radioisotope labels, for example), western blots, precipitation reactions, agglutination
assays (e.g., gel agglutination assays, hemagglutination assays), complement fixation
assays, immunofluorescence assays, protein A assays, and immunoelectrophoresis assays,
etc. In one embodiment, antibody binding is detected by detecting a label on the primary
antibody. In another embodiment, the primary antibody is detected by detecting binding

of a secondary antibody or reagent to the primary antibody. In a further embodiment, the secondary antibody is labeled. Many means are known in the art for detecting binding in an immunoassay and are within the scope of the present invention.

[0117] In another embodiment, where a ligand is identified, or the ability of a polypeptide fragment, variant or derivative of the invention to multimerize is being evaluated, binding can be assayed, e.g., by means well-known in the art, such as, for example, reducing and non-reducing gel chromatography, protein affinity chromatography, and affinity blotting. See generally, Phizicky et al., Microbiol. Rev. 59:94-123 (1995). In another embodiment, the ability of physiological correlates of a polypeptide of the present invention to bind to a substrate(s) of the polypeptide of the invention can be routinely assayed using techniques known in the art.

[0118] In addition, assays described herein (see Examples) and otherwise known in the art may routinely be applied to measure the ability of polypeptides of the present invention and fragments, variants and derivatives thereof to elicit polypeptide related biological activity (either *in vitro* or *in vivo*). Other methods will be known to the skilled artisan and are within the scope of the invention.

[0119]Of course, due to the degeneracy of the genetic code, one of ordinary skill in the art will immediately recognize that a large number of the nucleic acid molecules having a sequence at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99%, or 100% identical to, for example, the nucleic acid sequence of the cDNA contained in Clone ID NO:Z, a nucleic acid sequence referred to in Table 1A (e.g., SEQ ID NO:X), a nucleic acid sequence disclosed in Table 2 (e.g., the nucleic acid sequence delineated in columns 8 and 9) or fragments thereof, will encode polypeptides "having functional activity." In fact, since degenerate variants of any of these nucleotide sequences all encode the same polypeptide, in many instances, this will be clear to the skilled artisan even without performing the above described comparison assay. It will be further recognized in the art that, for such nucleic acid molecules that are not degenerate variants, a reasonable number will also encode a polypeptide having functional activity. This is because the skilled artisan is fully aware of amino acid substitutions that are either less likely or not likely to significantly effect protein function (e.g., replacing one aliphatic amino acid with a second aliphatic amino acid), as further described below.

[0120] For example, guidance concerning how to make phenotypically silent amino acid substitutions is provided in Bowie et al., "Deciphering the Message in Protein Sequences: Tolerance to Amino Acid Substitutions," Science 247:1306-1310 (1990), wherein the authors indicate that there are two main strategies for studying the tolerance of an amino acid sequence to change.

- [0121] The first strategy exploits the tolerance of amino acid substitutions by natural selection during the process of evolution. By comparing amino acid sequences in different species, conserved amino acids can be identified. These conserved amino acids are likely important for protein function. In contrast, the amino acid positions where substitutions have been tolerated by natural selection indicates that these positions are not critical for protein function. Thus, positions tolerating amino acid substitution could be modified while still maintaining biological activity of the protein.
- [0122] The second strategy uses genetic engineering to introduce amino acid changes at specific positions of a cloned gene to identify regions critical for protein function. For example, site directed mutagenesis or alanine-scanning mutagenesis (introduction of single alanine mutations at every residue in the molecule) can be used. See Cunningham et al., Science 244:1081-1085 (1989). The resulting mutant molecules can then be tested for biological activity.
- As the authors state, these two strategies have revealed that proteins are surprisingly tolerant of amino acid substitutions. The authors further indicate which amino acid changes are likely to be permissive at certain amino acid positions in the protein. For example, most buried (within the tertiary structure of the protein) amino acid residues require nonpolar side chains, whereas few features of surface side chains are generally conserved. Moreover, tolerated conservative amino acid substitutions involve replacement of the aliphatic or hydrophobic amino acids Ala, Val, Leu and Ile; replacement of the hydroxyl residues Ser and Thr; replacement of the acidic residues Asp and Glu; replacement of the amide residues Asn and Gln, replacement of the basic residues Lys, Arg, and His; replacement of the aromatic residues Phe, Tyr, and Trp, and replacement of the small-sized amino acids Ala, Ser, Thr, Met, and Gly. Besides conservative amino acid substitutions, variants of the present invention include (i) substitutions with one or more of the non-conserved amino acid residues, where the substituted amino acid residues may or may not be one encoded by the genetic code, or (ii)

substitutions with one or more of the amino acid residues having a substituent group, or (iii) fusion of the mature polypeptide with another compound, such as a compound to increase the stability and/or solubility of the polypeptide (for example, polyethylene glycol), or (iv) fusion of the polypeptide with additional amino acids, such as, for example, an IgG Fc fusion region peptide, serum albumin (preferably human serum albumin) or a fragment or variant thereof, or leader or secretory sequence, or a sequence facilitating purification. Such variant polypeptides are deemed to be within the scope of those skilled in the art from the teachings herein.

- [0124] For example, polypeptide variants containing amino acid substitutions of charged amino acids with other charged or neutral amino acids may produce proteins with improved characteristics, such as less aggregation. Aggregation of pharmaceutical formulations both reduces activity and increases clearance due to the aggregate's immunogenic activity. See Pinckard et al., Clin. Exp. Immunol. 2:331-340 (1967); Robbins et al., Diabetes 36: 838-845 (1987); Cleland et al., Crit. Rev. Therapeutic Drug Carrier Systems 10:307-377 (1993).
- the amino acid sequence of a polypeptide having an amino acid sequence which comprise the amino acid sequence of a polypeptide having an amino acid sequence which contains at least one amino acid substitution, but not more than 50 amino acid substitutions, even more preferably, not more than 40 amino acid substitutions, still more preferably, not more than 30 amino acid substitutions, and still even more preferably, not more than 20 amino acid substitutions from a polypeptide sequence disclosed herein. Of course it is highly preferable for a polypeptide to have an amino acid sequence which comprises the amino acid sequence of a polypeptide of SEQ ID NO:Y, an amino acid sequence encoded by SEQ ID NO:X, an amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, an amino acid sequence encoded by the complement of SEQ ID NO:X, and/or the amino acid sequence encoded by cDNA contained in Clone ID NO:Z which contains, in order of ever-increasing preference, at least one, but not more than 10, 9, 8, 7, 6, 5, 4, 3, 2 or 1 amino acid substitutions.
- [0126] In specific embodiments, the polypeptides of the invention comprise, or alternatively, consist of, fragments or variants of a reference amino acid sequence selected from: (a) the amino acid sequence of SEQ ID NO:Y or fragments thereof (e.g., the mature form and/or other fragments described herein); (b) the amino acid sequence encoded by

SEQ ID NO:X or fragments thereof; (c) the amino acid sequence encoded by the complement of SEQ ID NO:X or fragments thereof; (d) the amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or fragments thereof; and (e) the amino acid sequence encoded by cDNA contained in Clone ID NO:Z or fragments thereof; wherein the fragments or variants have 1-5, 5-10, 5-25, 5-50, 10-50 or 50-150, amino acid residue additions, substitutions, and/or deletions when compared to the reference amino acid sequence. In preferred embodiments, the amino acid substitutions are conservative. Polynucleotides encoding these polypeptides are also encompassed by the invention.

Polynucleotide and Polypeptide Fragments

The present invention is also directed to polynucleotide fragments of the [0127]polynucleotides (nucleic acids) of the invention. In the present invention, a "polynucleotide fragment" refers to a polynucleotide having a nucleic acid sequence which, for example: is a portion of the cDNA contained in Clone ID NO:Z or the complementary strand thereto; is a portion of the polynucleotide sequence encoding the polypeptide encoded by the cDNA contained in Clone ID NO:Z or the complementary strand thereto; is a portion of a polynucleotide sequence encoding the amino acid sequence encoded by the region of SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto; is a portion of the polynucleotide sequence of SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto; is a portion of the polynucleotide sequence in SEQ ID NO:X or the complementary strand thereto; is a polynucleotide sequence encoding a portion of the polypeptide of SEQ ID NO:Y; is a polynucleotide sequence encoding a portion of a polypeptide encoded by SEQ ID NO:X; is a polynucleotide sequence encoding a portion of a polypeptide encoded by the complement of the polynucleotide sequence in SEQ ID NO:X; is a portion of a polynucleotide sequence encoding the amino acid sequence encoded by the region of SEQ ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto; or is a portion of the polynucleotide sequence of SEQ ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto.

[0128] The polynucleotide fragments of the invention are preferably at least about 15 nt, and more preferably at least about 20 nt, still more preferably at least about 30 nt, and

even more preferably, at least about 40 nt, at least about 50 nt, at least about 75 nt, or at least about 150 nt in length. A fragment "at least 20 nt in length," for example, is intended to include 20 or more contiguous bases from the cDNA sequence contained in Clone ID NO:Z, or the nucleotide sequence shown in SEQ ID NO:X or the complementary stand thereto. In this context "about" includes the particularly recited value or a value larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. These nucleotide fragments have uses that include, but are not limited to, as diagnostic probes and primers as discussed herein. Of course, larger fragments (e.g., at least 160, 170, 180, 190, 200, 250, 500, 600, 1000, or 2000 nucleotides in length) are also encompassed by the invention.

Moreover, representative examples of polynucleotide fragments of the [0129] invention, comprise, or alternatively consist of, a sequence from about nucleotide number 1-50, 51-100, 101-150, 151-200, 201-250, 251-300, 301-350, 351-400, 401-450, 451-500, 501-550, 551-600, 651-700, 701-750, 751-800, 800-850, 851-900, 901-950, 951-1000, 1001-1050, 1051-1100, 1101-1150, 1151-1200, 1201-1250, 1251-1300, 1301-1350, 1351-1400, 1401-1450, 1451-1500, 1501-1550, 1551-1600, 1601-1650, 1651-1700, 1701-1750, 1751-1800, 1801-1850, 1851-1900, 1901-1950, 1951-2000, 2001-2050, 2051-2100, 2101-2150, 2151-2200, 2201-2250, 2251-2300, 2301-2350, 2351-2400, 2401-2450, 2451-2500, 2501-2550, 2551-2600, 2601-2650, 2651-2700, 2701-2750, 2751-2800, 2801-2850, 2851-2900, 2901-2950, 2951-3000, 3001-3050, 3051-3100, 3101-3150, 3151-3200, 3201-3250, 3251-3300, 3301-3350, 3351-3400, 3401-3450, 3451-3500, 3501-3550, 3551-3600, 3601-3650, 3651-3700, 3701-3750, 3751-3800, 3801-3850, 3851-3900, 3901-3950, 3951-4000, 4001-4050, 4051-4100, 4101-4150, 4151-4200, 4201-4250, 4251-4300, 4301-4350, 4351-4400, 4401-4450, 4451-4500, 4501-4550, 4551-4600, 4601-4650, 4651-4700, 4701-4750, 4751-4800, 4801-4850, 4851-4900, 4901-4950, 4951-5000, 5001-5050, 5051-5100, 5101-5150, 5151-5200, 5201-5250, 5251-5300, 5301-5350, 5351-5400, 5401-5450, 5451-5500, 5501-5550, 5551-5600, 5601-5650, 5651-5700, 5701-5750, 5751-5800, 5801-5850, 5851-5900, 5901-5950, 5951-6000, 6001-6050, 6051-6100, 6101-6150, 6151-6200, 6201-6250, 6251-6300, 6301-6350, 6351-6400, 6401-6450, 6451-6500, 6501-6550, 6551-6600, 6601-6650, 6651-6700, 6701-6750, 6751-6800, 6801-6850, 6851-6900, 6901-6950, 6951-7000, 7001-7050, 7051-7100, 7101-7150, 7151-7200, 7201-7250, 7251-7300 or 7301 to the end of SEQ ID NO:X, or the complementary strand thereto. In this context "about" includes

the particularly recited range or a range larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. Preferably, these fragments encode a polypeptide which has a functional activity (e.g., biological activity). More preferably, these polynucleotides can be used as probes or primers as discussed herein. Polynucleotides which hybridize to one or more of these polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

[0130] Further representative examples of polynucleotide fragments of the invention, comprise, or alternatively consist of, a sequence from about nucleotide number 1-50, 51-100, 101-150, 151-200, 201-250, 251-300, 301-350, 351-400, 401-450, 451-500, 501-550, 551-600, 651-700, 701-750, 751-800, 800-850, 851-900, 901-950, 951-1000, 1001-1050, 1051-1100, 1101-1150, 1151-1200, 1201-1250, 1251-1300, 1301-1350, 1351-1400, 1401-1450, 1451-1500, 1501-1550, 1551-1600, 1601-1650, 1651-1700, 1701-1750, 1751-1800, 1801-1850, 1851-1900, 1901-1950, 1951-2000, 2001-2050, 2051-2100, 2101-2150, 2151-2200, 2201-2250, 2251-2300, 2301-2350, 2351-2400, 2401-2450, 2451-2500, 2501-2550, 2551-2600, 2601-2650, 2651-2700, 2701-2750, 2751-2800, 2801-2850, 2851-2900, 2901-2950, 2951-3000, 3001-3050, 3051-3100, 3101-3150, 3151-3200, 3201-3250, 3251-3300, 3301-3350, 3351-3400, 3401-3450, 3451-3500, 3501-3550, 3551-3600, 3601-3650, 3651-3700, 3701-3750, 3751-3800, 3801-3850, 3851-3900, 3901-3950, 3951-4000, 4001-4050, 4051-4100, 4101-4150, 4151-4200, 4201-4250, 4251-4300, 4301-4350, 4351-4400, 4401-4450, 4451-4500, 4501-4550, 4551-4600, 4601-4650, 4651-4700, 4701-4750, 4751-4800, 4801-4850, 4851-4900, 4901-4950, 4951-5000, 5001-5050, 5051-5100, 5101-5150, 5151-5200, 5201-5250, 5251-5300, 5301-5350, 5351-5400, 5401-5450, 5451-5500, 5501-5550, 5551-5600, 5601-5650, 5651-5700, 5701-5750, 5751-5800, 5801-5850, 5851-5900, 5901-5950, 5951-6000, 6001-6050, 6051-6100, 6101-6150, 6151-6200, 6201-6250, 6251-6300, 6301-6350, 6351-6400, 6401-6450, 6451-6500, 6501-6550, 6551-6600, 6601-6650, 6651-6700, 6701-6750, 6751-6800, 6801-6850, 6851-6900, 6901-6950, 6951-7000, 7001-7050, 7051-7100, 7101-7150, 7151-7200, 7201-7250, 7251-7300 or 7301 to the end of the cDNA sequence contained in Clone ID NO:Z, or the complementary strand thereto. In this context "about" includes the particularly recited range or a range larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. Preferably, these fragments encode a polypeptide which has a functional activity (e.g., biological activity).

More preferably, these polynucleotides can be used as probes or primers as discussed herein. Polynucleotides which hybridize to one or more of these polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

[0131] Moreover, representative examples of polynucleotide fragments of the invention comprise, or alternatively consist of, a nucleic acid sequence comprising one, two, three, four, five, six, seven, eight, nine, ten, or more of the above described polynucleotide fragments of the invention in combination with a polynucleotide sequence delineated in Table 1B column 6. Additional, representative examples of polynucleotide fragments of the invention comprise, or alternatively consist of, a nucleic acid sequence comprising one, two, three, four, five, six, seven, eight, nine, ten, or more of the above described polynucleotide fragments of the invention in combination with a polynucleotide sequence that is the complementary strand of a sequence delineated in column 6 of Table 1B. In further embodiments, the above-described polynucleotide fragments of the invention comprise, or alternatively consist of, sequences delineated in Table 1B. column 6, and have a nucleic acid sequence which is different from that of the BAC fragment having the sequence disclosed in SEQ ID NO:B (see Table 1B, column 5). In additional embodiments, the above-described polynucleotide fragments of the invention comprise, or alternatively consist of, sequences delineated in Table 1B, column 6, and have a nucleic acid sequence which is different from that published for the BAC clone identified as BAC ID NO:A (see Table 1B, column 4). In additional embodiments, the above-described polynucleotides of the invention comprise, or alternatively consist of, sequences delineated Table 1B, column 6, and have a nucleic acid sequence which is different from that contained in the BAC clone identified as BAC ID NO:A (see Table 1B, column 4). Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides and polypeptides are also encompassed by the invention.

[0132] In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of, one, two, three, four, five, six, seven, eight, nine, ten, or more fragments of the sequences delineated in column 6 of Table 1B, and the polynucleotide sequence of SEQ ID NO:X (e.g., as defined in Table 1B, column 2) or fragments or

variants thereof. Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention.

- In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of, one, two, three, four, five, six, seven, eight, nine, ten, or more fragments of the sequences delineated in column 6 of Table 1B which correspond to the same Clone ID NO:Z (see Table 1B, column 1), and the polynucleotide sequence of SEQ ID NO:X (e.g., as defined in Table 1A or 1B) or fragments or variants thereof. Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention.
- In further specific embodiments, polynucleotides of the invention comprise, or alternatively consist of, one, two, three, four, five, six, seven, eight, nine, ten, or more fragments of the sequences delineated in the same row of column 6 of Table 1B, and the polynucleotide sequence of SEQ ID NO:X (e.g., as defined in Table 1A or 1B) or fragments or variants thereof. Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention.
- [0135] In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B and the 5' 10 polynucleotides of the sequence of SEQ ID NO:X are directly contiguous. Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.
- [0136] In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B and the 5' 10 polynucleotides of

a fragment or variant of the sequence of SEQ ID NO:X (e.g., as described herein) are directly contiguous Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids encoding these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.

In further specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of a fragment or variant of the sequence of SEQ ID NO:X and the 5' 10 polynucleotides of the sequence of one of the sequences delineated in column 6 of Table 1B are directly contiguous. Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids encoding these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.

In specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B and the 5' 10 polynucleotides of another sequence in column 6 are directly contiguous. In preferred embodiments, the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B is directly contiguous with the 5' 10 polynucleotides of the next sequential exon delineated in Table 1B, column 6. Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids encoding these polypeptides, and antibodies that bind these polypeptides are also

encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.

[0139] In the present invention, a "polypeptide fragment" refers to an amino acid sequence which is a portion of that contained in SEQ ID NO:Y, a portion of an amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, a portion of an amino acid sequence encoded by the polynucleotide sequence of SEO ID NO:X, a portion of an amino acid sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X, and/or a portion of an amino acid sequence encoded by the cDNA contained in Clone ID NO:Z. Protein (polypeptide) fragments may be "free-standing," or comprised within a larger polypeptide of which the fragment forms a part or region, most preferably as a single continuous region. Representative examples of polypeptide fragments of the invention, include, for example, fragments comprising, or alternatively consisting of, from about amino acid number 1-20, 21-40, 41-60, 61-80, 81-100, 102-120, 121-140, 141-160, 161-180, 181-200, 201-220, 221-240, 241-260, 261-280, 281-300, 301-320, 321-340, 341-360, 361-380, 381-400, 401-420, 421-440, 441-460, 461-480, 481-500, 501-520, 521-540, 541-560, 561-580, 581-600, 601-620, 621-640, 641-660, 661-680, 681-700, 701-720, 721-740, 741-760, 761-780, 781-800, 801-820, 821-840, 841-860, 861-880, 881-900, 901-920, 921-940, 941-960, 961-980, 981-1000, 1001-1020, 1021-1040, 1041-1060, 1061-1080, 1081-1100, 1101-1120, 1121-1140, 1141-1160, 1161-1180, 1181-1200, 1201-1220, 1221-1240, 1241-1260, 1261-1280, 1281-1300, 1301-1320, 1321-1340, 1341-1360, 1361-1380, 1381-1400, 1401-1420, 1421-1440, or 1441 to the end of the coding region. In a preferred embodiment, polypeptide fragments of the invention include, for example, fragments comprising, or alternatively consisting of, from about amino acid number 1-20, 21-40, 41-60, 61-80, 81-100, 102-120, 121-140, 141-160, 161-180, 181-200, 201-220, 221-240, 241-260, 261-280, 281-300, 301-320, 321-340, 341-360, 361-380, 381-400, 401-420, 421-440, 441-460, 461-480, 481-500, 501-520, 521-540, 541-560, 561-580, 581-600, 601-620, 621-640, 641-660, 661-680, 681-700, 701-720, 721-740, 741-760, 761-780, 781-800, 801-820, 821-840, 841-860, 861-880, 881-900, 901-920, 921-940, 941-960, 961-980, 981-1000, 1001-1020, 1021-1040, 1041-1060, 1061-1080, 1081-1100, 1101-1120, 1121-1140, 1141-1160, 1161-1180, 1181-1200, 1201-1220, 1221-1240, 1241-1260, 1261-1280, 1281-1300, 1301-1320, 1321-1340, 1341-1360, 1361-1380, 1381-

1400, 1401-1420, 1421-1440, or 1441 to the end of the coding region of SEQ ID NO:Y. Moreover, polypeptide fragments of the invention may be at least about 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 100, 110, 120, 130, 140, or 150 amino acids in length. In this context "about" includes the particularly recited ranges or values, or ranges or values larger or smaller by several (5, 4, 3, 2, or 1) amino acids, at either extreme or at both extremes. Polynucleotides encoding these polypeptide fragments are also encompassed by the invention.

- [0140] Even if deletion of one or more amino acids from the N-terminus of a protein results in modification of loss of one or more biological functions of the protein, other functional activities (e.g., biological activities, ability to multimerize, ability to bind a ligand) may still be retained. For example, the ability of shortened muteins to induce and/or bind to antibodies which recognize the complete or mature forms of the polypeptides generally will be retained when less than the majority of the residues of the complete or mature polypeptide are removed from the N-terminus. Whether a particular polypeptide lacking N-terminal residues of a complete polypeptide retains such immunologic activities can readily be determined by routine methods described herein and otherwise known in the art. It is not unlikely that a mutein with a large number of deleted N-terminal amino acid residues may retain some biological or immunogenic activities. In fact, peptides composed of as few as six amino acid residues may often evoke an immune response.
- [0141] Accordingly, polypeptide fragments include the secreted protein as well as the mature form. Further preferred polypeptide fragments include the secreted protein or the mature form having a continuous series of deleted residues from the amino or the carboxy terminus, or both. For example, any number of amino acids, ranging from 1-60, can be deleted from the amino terminus of either the secreted polypeptide or the mature form. Similarly, any number of amino acids, ranging from 1-30, can be deleted from the carboxy terminus of the secreted protein or mature form. Furthermore, any combination of the above amino and carboxy terminus deletions is preferred. Similarly, polynucleotides encoding these polypeptide fragments are also preferred.
- [0142] The present invention further provides polypeptides having one or more residues deleted from the amino terminus of the amino acid sequence of a polypeptide disclosed herein (e.g., a polypeptide of SEQ ID NO:Y, a polypeptide encoded by the

polynucleotide sequence contained in SEQ ID NO:X or the complement thereof, a polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, a polypeptide encoded by the portion of SEQ ID NO:B as defined in column 6 of Table 1B, and/or a polypeptide encoded by the cDNA contained in Clone ID NO:Z). In particular, N-terminal deletions may be described by the general formula m-q, where q is a whole integer representing the total number of amino acid residues in a polypeptide of the invention (e.g., the polypeptide disclosed in SEQ ID NO:Y, or the polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2), and m is defined as any integer ranging from 2 to q-6. Polynucleotides encoding these polypeptides are also encompassed by the invention.

- The present invention further provides polypeptides having one or more residues from the carboxy terminus of the amino acid sequence of a polypeptide disclosed herein (e.g., a polypeptide of SEQ ID NO:Y, a polypeptide encoded by the polynucleotide sequence contained in SEQ ID NO:X, a polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, and/or a polypeptide encoded by the cDNA contained in Clone ID NO:Z). In particular, C-terminal deletions may be described by the general formula 1-n, where n is any whole integer ranging from 6 to q-1, and where n corresponds to the position of amino acid residue in a polypeptide of the invention. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- In addition, any of the above described N- or C-terminal deletions can be combined to produce a N- and C-terminal deleted polypeptide. The invention also provides polypeptides having one or more amino acids deleted from both the amino and the carboxyl termini, which may be described generally as having residues m-n of a polypeptide encoded by SEQ ID NO:X (e.g., including, but not limited to, the preferred polypeptide disclosed as SEQ ID NO:Y and the polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2), the cDNA contained in Clone ID NO:Z, and/or the complement thereof, where n and m are integers as described above. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- [0145] Also as mentioned above, even if deletion of one or more amino acids from the C-terminus of a protein results in modification of loss of one or more biological functions of the protein, other functional activities (e.g., biological activities, ability to multimerize, ability to bind a ligand) may still be retained. For example the ability of the shortened

mutein to induce and/or bind to antibodies which recognize the complete or mature forms of the polypeptide generally will be retained when less than the majority of the residues of the complete or mature polypeptide are removed from the C-terminus. Whether a particular polypeptide lacking C-terminal residues of a complete polypeptide retains such immunologic activities can readily be determined by routine methods described herein and otherwise known in the art. It is not unlikely that a mutein with a large number of deleted C-terminal amino acid residues may retain some biological or immunogenic activities. In fact, peptides composed of as few as six amino acid residues may often evoke an immune response.

- [0146] The present application is also directed to proteins containing polypeptides at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a polypeptide sequence set forth herein. In preferred embodiments, the application is directed to proteins containing polypeptides at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to polypeptides having the amino acid sequence of the specific N- and C-terminal deletions. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- [0147] Any polypeptide sequence encoded by, for example, the polynucleotide sequences set forth as SEQ ID NO:X or the complement thereof, (presented, for example, in Tables 1A and 2), the cDNA contained in Clone ID NO:Z, or the polynucleotide sequence as defined in column 6 of Table 1B, may be analyzed to determine certain preferred regions of the polypeptide. For example, the amino acid sequence of a polypeptide encoded by a polynucleotide sequence of SEQ ID NO:X (e.g., the polypeptide of SEQ ID NO:Y and the polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2) or the cDNA contained in Clone ID NO:Z may be analyzed using the default parameters of the DNASTAR computer algorithm (DNASTAR, Inc., 1228 S. Park St., Madison, WI 53715 USA; http://www.dnastar.com/).
- [0148] Polypeptide regions that may be routinely obtained using the DNASTAR computer algorithm include, but are not limited to, Garnier-Robson alpha-regions, beta-regions, turn-regions, and coil-regions; Chou-Fasman alpha-regions, beta-regions, and turn-regions; Kyte-Doolittle hydrophilic regions and hydrophobic regions; Eisenberg alpha- and beta-amphipathic regions; Karplus-Schulz flexible regions; Emini surface-forming regions; and Jameson-Wolf regions of high antigenic index. Among

highly preferred polynucleotides of the invention in this regard are those that encode polypeptides comprising regions that combine several structural features, such as several (e.g., 1, 2, 3 or 4) of the features set out above.

- [0149] Additionally, Kyte-Doolittle hydrophilic regions and hydrophobic regions, Emini surface-forming regions, and Jameson-Wolf regions of high antigenic index (i.e., containing four or more contiguous amino acids having an antigenic index of greater than or equal to 1.5, as identified using the default parameters of the Jameson-Wolf program) can routinely be used to determine polypeptide regions that exhibit a high degree of potential for antigenicity. Regions of high antigenicity are determined from data by DNASTAR analysis by choosing values which represent regions of the polypeptide which are likely to be exposed on the surface of the polypeptide in an environment in which antigen recognition may occur in the process of initiation of an immune response.
- [0150] Preferred polypeptide fragments of the invention are fragments comprising, or alternatively, consisting of, an amino acid sequence that displays a functional activity (e.g. biological activity) of the polypeptide sequence of which the amino acid sequence is a fragment. By a polypeptide displaying a "functional activity" is meant a polypeptide capable of one or more known functional activities associated with a full-length protein, such as, for example, biological activity, antigenicity, immunogenicity, and/or multimerization, as described herein.
- [0151] Other preferred polypeptide fragments are biologically active fragments. Biologically active fragments are those exhibiting activity similar, but not necessarily identical, to an activity of the polypeptide of the present invention. The biological activity of the fragments may include an improved desired activity, or a decreased undesirable activity.
- [0152] In preferred embodiments, polypeptides of the invention comprise, or alternatively consist of, one, two, three, four, five or more of the antigenic fragments of the polypeptide of SEQ ID NO:Y, or portions thereof. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- [0153] The present invention encompasses polypeptides comprising, or alternatively consisting of, an epitope of: the polypeptide sequence shown in SEQ ID NO:Y; a polypeptide sequence encoded by SEQ ID NO:X or the complementary strand thereto; the polypeptide sequence encoded by the portion of SEQ ID NO:X as defined in columns 8

and 9 of Table 2; the polypeptide sequence encoded by the portion of SEQ ID NO:B as defined in column 6 of Table 1B or the complement thereto; the polypeptide sequence encoded by the cDNA contained in Clone ID NO:Z; or the polypeptide sequence encoded by a polynucleotide that hybridizes to the sequence of SEQ ID NO:X, the complement of the sequence of SEQ ID NO:X, the complement of a portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, or the cDNA sequence contained in Clone ID NO:Z under stringent hybridization conditions or alternatively, under lower stringency hybridization as defined *supra*. The present invention further encompasses polynucleotide sequences encoding an epitope of a polypeptide sequence of the invention (such as, for example, the sequence disclosed in SEQ ID NO:X, or a fragment thereof), polynucleotide sequences of the complementary strand of a polynucleotide sequence encoding an epitope of the invention, and polynucleotide sequences which hybridize to the complementary strand under stringent hybridization conditions or alternatively, under lower stringency hybridization conditions defined *supra*.

In a preferred embodiment, the present invention encompasses a polypeptide comprising an epitope, as well as the polynucleotide encoding this polypeptide. An "immunogenic epitope," as used herein, is defined as a portion of a protein that elicits an antibody response in an animal, as determined by any method known in the art, for example, by the methods for generating antibodies described *infra*. (See, for example, Geysen et al., Proc. Natl. Acad. Sci. USA 81:3998- 4002 (1983)). The term "antigenic epitope," as used herein, is defined as a portion of a protein to which an antibody can immunospecifically bind its antigen as determined by any method well known in the art, for example, by the immunoassays described herein. Immunospecific binding excludes non-specific binding but does not necessarily exclude cross- reactivity with other antigens. Antigenic epitopes need not necessarily be immunogenic.

[0155] Fragments which function as epitopes may be produced by any conventional means. (See, e.g., Houghten, R. A., Proc. Natl. Acad. Sci. USA 82:5131-5135 (1985) further described in U.S. Patent No. 4,631,211.)

[0156] In the present invention, antigenic epitopes preferably contain a sequence of at least 4, at least 5, at least 6, at least 7, more preferably at least 8, at least 9, at least 10, at

least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 25, at least 30, at least 40, at least 50, and, most preferably, between about 15 to about 30 amino acids. Preferred polypeptides comprising immunogenic or antigenic epitopes are at least 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, or 100 amino acid residues in length. Additional non-exclusive preferred antigenic epitopes include the antigenic epitopes disclosed herein, as well as portions thereof. Antigenic epitopes are useful, for example, to raise antibodies, including monoclonal antibodies, that specifically bind the epitope. Preferred antigenic epitopes include the antigenic epitopes disclosed herein, as well as any combination of two, three, four, five or more of these antigenic epitopes. Antigenic epitopes can be used as the target molecules in immunoassays. (See, for instance, Wilson et al., Cell 37:767-778 (1984); Sutcliffe et al., Science 219:660-666 (1983)).

Non-limiting examples of epitopes of polypeptides that can be used to generate [0157] antibodies of the invention include a polypeptide comprising, or alternatively consisting of, at least one, two, three, four, five, six or more of the portion(s) of SEQ ID NO:Y specified in column 6 of Table 1A. These polypeptide fragments have been determined to bear antigenic epitopes of the proteins of the invention by the analysis of the Jameson-Wolf antigenic index which is included in the DNAStar suite of computer programs. By "comprise" it is intended that a polypeptide contains at least one, two, three, four, five, six or more of the portion(s) of SEQ ID NO:Y shown in column 6 of Table 1A, but it may contain additional flanking residues on either the amino or carboxyl termini of the recited portion. Such additional flanking sequences are preferably sequences naturally found adjacent to the portion; i.e., contiguous sequence shown in SEQ ID NO:Y. The flanking sequence may, however, be sequences from a heterologous polypeptide, such as from another protein described herein or from a heterologous polypeptide not described herein. In particular embodiments, epitope portions of a polypeptide of the invention comprise one, two, three, or more of the portions of SEQ ID NO:Y shown in column 6 of Table 1A. Polynucleotides encoding these polypeptides are also encompassed by the invention.

[0158] Similarly, immunogenic epitopes can be used, for example, to induce antibodies according to methods well known in the art. See, for instance, Sutcliffe et al., supra; Wilson et al., supra; Chow et al., Proc. Natl. Acad. Sci. USA 82:910-914; and Bittle et al.; J. Gen. Virol. 66:2347-2354 (1985). Preferred immunogenic epitopes

include the immunogenic epitopes disclosed herein, as well as any combination of two, three, four, five or more of these immunogenic epitopes. The polypeptides comprising one or more immunogenic epitopes may be presented for eliciting an antibody response together with a carrier protein, such as an albumin, to an animal system (such as rabbit or mouse), or, if the polypeptide is of sufficient length (at least about 25 amino acids), the polypeptide may be presented without a carrier. However, immunogenic epitopes comprising as few as 8 to 10 amino acids have been shown to be sufficient to raise antibodies capable of binding to, at the very least, linear epitopes in a denatured polypeptide (e.g., in Western blotting).

[0159]Epitope-bearing polypeptides of the present invention may be used to induce antibodies according to methods well known in the art including, but not limited to, in vivo immunization, in vitro immunization, and phage display methods. See, e.g., Sutcliffe et al., supra; Wilson et al., supra, and Bittle et al., J. Gen. Virol., 66:2347-2354 (1985). If in vivo immunization is used, animals may be immunized with free peptide; however, antipeptide antibody titer may be boosted by coupling the peptide to a macromolecular carrier. such as keyhole limpet hemacyanin (KLH) or tetanus toxoid. For instance, peptides containing cysteine residues may be coupled to a carrier using a linker such as maleimidobenzoyl- N-hydroxysuccinimide ester (MBS), while other peptides may be coupled to carriers using a more general linking agent such as glutaraldehyde. Animals such as rabbits, rats and mice are immunized with either free or carrier- coupled peptides, for instance, by intraperitoneal and/or intradermal injection of emulsions containing about 100 μg of peptide or carrier protein and Freund's adjuvant or any other adjuvant known for stimulating an immune response. Several booster injections may be needed, for instance, at intervals of about two weeks, to provide a useful titer of anti-peptide antibody which can be detected, for example, by ELISA assay using free peptide adsorbed to a solid surface. The titer of anti-peptide antibodies in serum from an immunized animal may be increased by selection of anti-peptide antibodies, for instance, by adsorption to the peptide on a solid support and elution of the selected antibodies according to methods well known in the art.

[0160] As one of skill in the art will appreciate, and as discussed above, the polypeptides of the present invention (e.g., those comprising an immunogenic or antigenic epitope) can be fused to heterologous polypeptide sequences. For example, polypeptides of the present

invention (including fragments or variants thereof), may be fused with the constant domain of immunoglobulins (IgA, IgE, IgG, IgM), or portions thereof (CH1, CH2, CH3, or any combination thereof and portions thereof, resulting in chimeric polypeptides. By way of another non-limiting example, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) may be fused with albumin (including but not limited to recombinant human serum albumin or fragments or variants thereof (see, e.g., U.S. Patent No. 5,876,969, issued March 2, 1999, EP Patent 0 413 622, and U.S. Patent No. 5,766,883, issued June 16, 1998, herein incorporated by reference in their entirety)). In a preferred embodiment, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) are fused with the mature form of human serum albumin (i.e., amino acids 1 - 585 of human serum albumin as shown in Figures 1 and 2 of EP Patent 0 322 094) which is herein incorporated by reference in its entirety. In another preferred embodiment, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) are fused with polypeptide fragments comprising, or alternatively consisting of, amino acid residues 1-z of human serum albumin, where z is an integer from 369 to 419, as described in U.S. Patent 5,766,883 herein incorporated by reference in its entirety. Polypeptides and/or antibodies of the present invention (including fragments or variants thereof) may be fused to either the Nor C-terminal end of the heterologous protein (e.g., immunoglobulin Fc polypeptide or human serum albumin polypeptide). Polynucleotides encoding fusion proteins of the invention are also encompassed by the invention.

[0161] Such fusion proteins as those described above may facilitate purification and may increase half-life *in vivo*. This has been shown for chimeric proteins consisting of the first two domains of the human CD4-polypeptide and various domains of the constant regions of the heavy or light chains of mammalian immunoglobulins. See, e.g., EP 394,827; Traunecker et al., Nature, 331:84-86 (1988). Enhanced delivery of an antigen across the epithelial barrier to the immune system has been demonstrated for antigens (e.g., insulin) conjugated to an FcRn binding partner such as IgG or Fc fragments (see, e.g., PCT Publications WO 96/22024 and WO 99/04813). IgG Fusion proteins that have a disulfide-linked dimeric structure due to the IgG portion desulfide bonds have also been found to be more efficient in binding and neutralizing other molecules than monomeric polypeptides or fragments thereof alone. See, e.g., Fountoulakis et al., J. Biochem.,

270:3958-3964 (1995). Nucleic acids encoding the above epitopes can also be recombined with a gene of interest as an epitope tag (e.g., the hemagglutinin (HA) tag or flag tag) to aid in detection and purification of the expressed polypeptide. For example, a system described by Janknecht et al. allows for the ready purification of non-denatured fusion proteins expressed in human cell lines (Janknecht et al., 1991, Proc. Natl. Acad. Sci. USA 88:8972-897). In this system, the gene of interest is subcloned into a vaccinia recombination plasmid such that the open reading frame of the gene is translationally fused to an amino-terminal tag consisting of six histidine residues. The tag serves as a matrix binding domain for the fusion protein. Extracts from cells infected with the recombinant vaccinia virus are loaded onto Ni2+ nitriloacetic acid-agarose column and histidine-tagged proteins can be selectively eluted with imidazole-containing buffers.

Fusion Proteins

- [0162] Any polypeptide of the present invention can be used to generate fusion proteins. For example, the polypeptide of the present invention, when fused to a second protein, can be used as an antigenic tag. Antibodies raised against the polypeptide of the present invention can be used to indirectly detect the second protein by binding to the polypeptide. Moreover, because secreted proteins target cellular locations based on trafficking signals, polypeptides of the present invention which are shown to be secreted can be used as targeting molecules once fused to other proteins.
- [0163] Examples of domains that can be fused to polypeptides of the present invention include not only heterologous signal sequences, but also other heterologous functional regions. The fusion does not necessarily need to be direct, but may occur through linker sequences.
- [0164] In certain preferred embodiments, proteins of the invention are fusion proteins comprising an amino acid sequence that is an N and/or C- terminal deletion of a polypeptide of the invention. In preferred embodiments, the invention is directed to a fusion protein comprising an amino acid sequence that is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a polypeptide sequence of the invention. Polynucleotides encoding these proteins are also encompassed by the invention.
- [0165] Moreover, fusion proteins may also be engineered to improve characteristics of the polypeptide of the present invention. For instance, a region of additional amino acids.

particularly charged amino acids, may be added to the N-terminus of the polypeptide to improve stability and persistence during purification from the host cell or subsequent handling and storage. Also, peptide moieties may be added to the polypeptide to facilitate purification. Such regions may be removed prior to final preparation of the polypeptide. The addition of peptide moieties to facilitate handling of polypeptides are familiar and routine techniques in the art.

[0166] As one of skill in the art will appreciate that, as discussed above, polypeptides of the present invention, and epitope-bearing fragments thereof, can be combined with heterologous polypeptide sequences. For example, the polypeptides of the present invention may be fused with heterologous polypeptide sequences, for example, the polypeptides of the present invention may be fused with the constant domain of immunoglobulins (IgA, IgE, IgG, IgM) or portions thereof (CH1, CH2, CH3, and any combination thereof, including both entire domains and portions thereof), or albumin (including, but not limited to, native or recombinant human albumin or fragments or variants thereof (see, e.g., U.S. Patent No. 5,876,969, issued March 2, 1999, EP Patent 0 413 622, and U.S. Patent No. 5,766,883, issued June 16, 1998, herein incorporated by reference in their entirety)), resulting in chimeric polypeptides. For example, EP-A-O 464 533 (Canadian counterpart 2045869) discloses fusion proteins comprising various portions of constant region of immunoglobulin molecules together with another human protein or part thereof. In many cases, the Fc part in a fusion protein is beneficial in therapy and diagnosis, and thus can result in, for example, improved pharmacokinetic properties (EP-A 0232 262). Alternatively, deleting the Fc part after the fusion protein has been expressed, detected, and purified, would be desired. For example, the Fc portion may hinder therapy and diagnosis if the fusion protein is used as an antigen for immunizations. In drug discovery, for example, human proteins, such as hIL-5, have been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists of hIL-5. See, D. Bennett et al., J. Molecular Recognition 8:52-58 (1995); K. Johanson et al., J. Biol. Chem. 270:9459-9471 (1995).

[0167] Moreover, the polypeptides of the present invention can be fused to marker sequences, such as a polypeptide which facilitates purification of the fused polypeptide. In preferred embodiments, the marker amino acid sequence is a hexa-histidine peptide, such as the tag provided in a pQE vector (QIAGEN, Inc., 9259 Eton Avenue, Chatsworth,

CA, 91311), among others, many of which are commercially available. As described in Gentz et al., Proc. Natl. Acad. Sci. USA 86:821-824 (1989), for instance, hexa-histidine provides for convenient purification of the fusion protein. Another peptide tag useful for purification, the "HA" tag, corresponds to an epitope derived from the influenza hemagglutinin protein (Wilson et al., Cell 37:767 (1984).)

Additional fusion proteins of the invention may be generated through the [0168]techniques of gene-shuffling, motif-shuffling, exon-shuffling, and/or codon-shuffling (collectively referred to as "DNA shuffling"), briefly described below, and further described herein. DNA shuffling may be employed to modulate the activities of polypeptides of the invention, such methods can be used to generate polypeptides with altered activity, as well as agonists and antagonists of the polypeptides. See, generally, U.S. Patent Nos. 5,605,793; 5,811,238; 5,830,721; 5,834,252; and 5,837,458, and Patten et al., Curr. Opinion Biotechnol. 8:724-33 (1997); Harayama, Trends Biotechnol. 16(2):76-82 (1998); Hansson et al., J. Mol. Biol. 287:265-76 (1999); and Lorenzo and Blasco, Biotechniques 24(2):308-13 (1998); each of these patents and publications are hereby incorporated by reference in its entirety). In a preferred embodiment, one or more components, motifs, sections, parts, domains, fragments, etc., of a polynucleotide encoding a polypeptide of the invention may be recombined with one or more components, motifs, sections, parts, domains, fragments, etc., of one or more heterologous molecules encoding a heterologous polypeptide.

[0169] Thus, any of these above fusions can be engineered using the polynucleotides or the polypeptides of the present invention.

Recombinant and Synthetic Production of Polypeptides of the Invention

[0170] The present invention also relates to vectors containing the polynucleotide of the present invention, host cells, and the production of polypeptides by synthetic and recombinant techniques. The vector may be, for example, a phage, plasmid, viral, or retroviral vector. Retroviral vectors may be replication competent or replication defective. In the latter case, viral propagation generally will occur only in complementing host cells.

[0171] The polynucleotides of the invention may be joined to a vector containing a selectable marker for propagation in a host. Generally, a plasmid vector is introduced in a precipitate, such as a calcium phosphate precipitate, or in a complex with a charged lipid.

If the vector is a virus, it may be packaged *in vitro* using an appropriate packaging cell line and then transduced into host cells.

[0172] The polynucleotide insert should be operatively linked to an appropriate promoter, such as the phage lambda PL promoter, the E. coli lac, trp, phoA and tac promoters, the SV40 early and late promoters and promoters of retroviral LTRs, to name a few. Other suitable promoters will be known to the skilled artisan. The expression constructs will further contain sites for transcription initiation, termination, and, in the transcribed region, a ribosome binding site for translation. The coding portion of the transcripts expressed by the constructs will preferably include a translation initiating codon at the beginning and a termination codon (UAA, UGA or UAG) appropriately positioned at the end of the polypeptide to be translated.

[0173] As indicated, the expression vectors will preferably include at least one selectable marker. Such markers include dihydrofolate reductase, G418 or neomycin resistance, glutamine synthase, for eukaryotic cell culture and tetracycline, kanamycin or ampicillin resistance genes for culturing in *E. coli* and other bacteria. Representative examples of appropriate hosts include, but are not limited to, bacterial cells, such as *E. coli*, Streptomyces and Salmonella typhimurium cells; fungal cells, such as yeast cells (e.g., Saccharomyces cerevisiae or Pichia pastoris (ATCC Accession No. 201178)); insect cells such as Drosophila S2 and Spodoptera Sf9 cells; animal cells such as CHO, COS, 293, NSO and Bowes melanoma cells; and plant cells. Appropriate culture mediums and conditions for the above-described host cells are known in the art.

[0174] Among vectors preferred for use in bacteria include pQE70, pQE60 and pQE-9, available from QIAGEN, Inc.; pBluescript vectors, Phagescript vectors, pNH8A, pNH16a, pNH18A, pNH46A, available from Stratagene Cloning Systems, Inc.; and ptrc99a, pKK223-3, pKK233-3, pDR540, pRIT5 available from Pharmacia Biotech, Inc. Among preferred eukaryotic vectors are pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; and pSVK3, pBPV, pMSG and pSVL available from Pharmacia. Preferred expression vectors for use in yeast systems include, but are not limited to pYES2, pYD1, pTEF1/Zeo, pYES2/GS, pPICZ, pGAPZ, pGAPZalph, pPIC9, pPIC3.5, pHIL-D2, pHIL-S1, pPIC3.5K, pPIC9K, and PAO815 (all available from Invitrogen, Carlsbad, CA). Other suitable vectors will be readily apparent to the skilled artisan.

[0175]Vectors which use glutamine synthase (GS) or DHFR as the selectable markers can be amplified in the presence of the drugs methionine sulphoximine or methotrexate, respectively. An advantage of glutamine synthase based vectors are the availabilty of cell lines (e.g., the murine myeloma cell line, NS0) which are glutamine synthase negative. Glutamine synthase expression systems can also function in glutamine synthase expressing cells (e.g., Chinese Hamster Ovary (CHO) cells) by providing additional inhibitor to prevent the functioning of the endogenous gene. A glutamine synthase expression system and components thereof are detailed in PCT publications: WO87/04462; WO86/05807; WO89/01036; WO89/10404; and WO91/06657 which are hereby incorporated in their entireties by reference herein. Additionally, glutamine synthase expression vectors can be obtained from Lonza Biologics, Inc. (Portsmouth, NH). Expression and production of monoclonal antibodies using a GS expression system in murine myeloma cells is described in Bebbington et al., Bio/technology 10:169(1992) and in Biblia and Robinson Biotechnol. Prog. 11:1 (1995) which are herein incorporated by reference.

Vector constructs described herein, and additionally encompasses host cells containing nucleotide sequences of the invention that are operably associated with one or more heterologous control regions (e.g., promoter and/or enhancer) using techniques known of in the art. The host cell can be a higher eukaryotic cell, such as a mammalian cell (e.g., a human derived cell), or a lower eukaryotic cell, such as a yeast cell, or the host cell can be a prokaryotic cell, such as a bacterial cell. A host strain may be chosen which modulates the expression of the inserted gene sequences, or modifies and processes the gene product in the specific fashion desired. Expression from certain promoters can be elevated in the presence of certain inducers; thus expression of the genetically engineered polypeptide may be controlled. Furthermore, different host cells have characteristics and specific mechanisms for the translational and post-translational processing and modification (e.g., phosphorylation, cleavage) of proteins. Appropriate cell lines can be chosen to ensure the desired modifications and processing of the foreign protein expressed.

[0177] Introduction of the nucleic acids and nucleic acid constructs of the invention into the host cell can be effected by calcium phosphate transfection, DEAE-dextran mediated transfection, cationic lipid-mediated transfection, electroporation, transduction,

infection, or other methods. Such methods are described in many standard laboratory manuals, such as Davis et al., Basic Methods In Molecular Biology (1986). It is specifically contemplated that the polypeptides of the present invention may in fact be expressed by a host cell lacking a recombinant vector.

In addition to encompassing host cells containing the vector constructs discussed [0178] herein, the invention also encompasses primary, secondary, and immortalized host cells of vertebrate origin, particularly mammalian origin, that have been engineered to delete or replace endogenous genetic material (e.g., immune/hematopoietic antigen coding sequence), and/or to include genetic material (e.g., heterologous polynucleotide sequences) that is operably associated with immune/hematopoietic associated polynucleotides of the invention, and which activates, alters, and/or amplifies endogenous immune/hematopoietic associated polynucleotides. For example, techniques known in the art may be used to operably associate heterologous control regions (e.g., promoter and/or enhancer) and endogenous immune/hematopoietic associated polynucleotide sequences via homologous recombination (see, e.g., U.S. Patent Number 5,641,670, issued June 24, 1997; International Publication Number WO 96/29411; International Publication Number WO 94/12650; Koller et al., Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); and Zijlstra et al., Nature 342:435-438 (1989), the disclosures of each of which are incorporated by reference in their entireties).

Polypeptides of the present invention can also be recovered from: products purified from natural sources, including bodily fluids, tissues and cells, whether directly isolated or cultured; products of chemical synthetic procedures; and products produced by recombinant techniques from a prokaryotic or eukaryotic host, including, for example, bacterial, yeast, higher plant, insect, and mammalian cells. Depending upon the host employed in a recombinant production procedure, the polypeptides of the present invention may be glycosylated or may be non-glycosylated. In addition, polypeptides of the invention may also include an initial modified methionine residue, in some cases as a result of host-mediated processes. Thus, it is well known in the art that the N-terminal methionine encoded by the translation initiation codon generally is removed with high efficiency from any protein after translation in all eukaryotic cells. While the N-terminal methionine on most proteins also is efficiently removed in most prokaryotes, for some

proteins, this prokaryotic removal process is inefficient, depending on the nature of the amino acid to which the N-terminal methionine is covalently linked.

In one embodiment, the yeast Pichia pastoris is used to express polypeptides of [0180] the invention in a eukaryotic system. Pichia pastoris is a methylotrophic yeast which can metabolize methanol as its sole carbon source. A main step in the methanol metabolization pathway is the oxidation of methanol to formaldehyde using O2. This reaction is catalyzed by the enzyme alcohol oxidase. In order to metabolize methanol as its sole carbon source, Pichia pastoris must generate high levels of alcohol oxidase due, in part, to the relatively low affinity of alcohol oxidase for O2. Consequently, in a growth medium depending on methanol as a main carbon source, the promoter region of one of the two alcohol oxidase genes (AOXI) is highly active. In the presence of methanol, alcohol oxidase produced from the AOXI gene comprises up to approximately 30% of the total soluble protein in Pichia pastoris. See, Ellis, S.B., et al., Mol. Cell. Biol. 5:1111-21 (1985); Koutz, P.J, et al., Yeast 5:167-77 (1989); Tschopp, J.F., et al., Nucl. Acids Res. 15:3859-76 (1987). Thus, a heterologous coding sequence, such as, for example, a polynucleotide of the present invention, under the transcriptional regulation of all or part of the AOXI regulatory sequence is expressed at exceptionally high levels in Pichia yeast grown in the presence of methanol.

In one example, the plasmid vector pPIC9K is used to express DNA encoding a polypeptide of the invention, as set forth herein, in a *Pichea* yeast system essentially as described in "*Pichia* Protocols: Methods in Molecular Biology," D.R. Higgins and J. Cregg, eds. The Humana Press, Totowa, NJ, 1998. This expression vector allows expression and secretion of a polypeptide of the invention by virtue of the strong *AOXI* promoter linked to the *Pichia pastoris* alkaline phosphatase (PHO) secretory signal peptide (i.e., leader) located upstream of a multiple cloning site.

[0182] Many other yeast vectors could be used in place of pPIC9K, such as, pYES2, pYD1, pTEF1/Zeo, pYES2/GS, pPICZ, pGAPZ, pGAPZalpha, pPIC9, pPIC3.5, pHIL-D2, pHIL-S1, pPIC3.5K, and PAO815, as one skilled in the art would readily appreciate, as long as the proposed expression construct provides appropriately located signals for

transcription, translation, secretion (if desired), and the like, including an in-frame AUG as required.

[0183] In another embodiment, high-level expression of a heterologous coding sequence, such as, for example, a polynucleotide of the present invention, may be achieved by cloning the heterologous polynucleotide of the invention into an expression vector such as, for example, pGAPZ or pGAPZalpha, and growing the yeast culture in the absence of methanol.

[0184] In addition to encompassing host cells containing the vector constructs discussed herein, the invention also encompasses primary, secondary, and immortalized host cells of vertebrate origin, particularly mammalian origin, that have been engineered to delete or replace endogenous genetic material (e.g., coding sequence), and/or to include genetic material (e.g., heterologous polynucleotide sequences) that is operably associated with polynucleotides of the invention, and which activates, alters, and/or amplifies endogenous polynucleotides. For example, techniques known in the art may be used to operably associate heterologous control regions (e.g., promoter and/or enhancer) and endogenous polynucleotide sequences via homologous recombination (see, e.g., U.S. Patent No. 5,641,670, issued June 24, 1997; International Publication No. WO 96/29411, published September 26, 1996; International Publication No. WO 94/12650, published August 4, 1994; Koller et al., Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); and Zijlstra et al., Nature 342:435-438 (1989), the disclosures of each of which are incorporated by reference in their entireties).

[0185] In addition, polypeptides of the invention can be chemically synthesized using techniques known in the art (e.g., see Creighton, 1983, Proteins: Structures and Molecular Principles, W.H. Freeman & Co., N.Y., and Hunkapiller et al., Nature, 310:105-111 (1984)). For example, a polypeptide corresponding to a fragment of a polypeptide can be synthesized by use of a peptide synthesizer. Furthermore, if desired, nonclassical amino acids or chemical amino acid analogs can be introduced as a substitution or addition into the polypeptide sequence. Non-classical amino acids include, but are not limited to, to the D-isomers of the common amino acids, 2,4-diaminobutyric acid, a-amino isobutyric acid, 4-aminobutyric acid, Abu, 2-amino butyric acid, g-Abu, e-Ahx, 6-amino hexanoic acid, Aib, 2-amino isobutyric acid, 3-amino propionic acid, ornithine, norleucine, norvaline, hydroxyproline, sarcosine, citrulline, homocitrulline, cysteic acid, t-butylglycine, t-

butylalanine, phenylglycine, cyclohexylalanine, b-alanine, fluoro-amino acids, designer amino acids such as b-methyl amino acids, Ca-methyl amino acids, Na-methyl amino acids, and amino acid analogs in general. Furthermore, the amino acid can be D (dextrorotary) or L (levorotary).

[0186] The invention encompasses polypeptides of the present invention which are differentially modified during or after translation, e.g., by glycosylation, acetylation, phosphorylation, amidation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to an antibody molecule or other cellular ligand, etc. Any of numerous chemical modifications may be carried out by known techniques, including but not limited, to specific chemical cleavage by cyanogen bromide, trypsin, chymotrypsin, papain, V8 protease, NaBH₄; acetylation, formylation, oxidation, reduction; metabolic synthesis in the presence of tunicamycin; etc.

[0187] Additional post-translational modifications encompassed by the invention include, for example, e.g., N-linked or O-linked carbohydrate chains, processing of N-terminal or C-terminal ends), attachment of chemical moieties to the amino acid backbone, chemical modifications of N-linked or O-linked carbohydrate chains, and addition or deletion of an N-terminal methionine residue as a result of procaryotic host cell expression. The polypeptides may also be modified with a detectable label, such as an enzymatic, fluorescent, isotopic or affinity label to allow for detection and isolation of the protein.

[0188] Examples of suitable enzymes include horseradish peroxidase, alkaline phosphatase, beta-galactosidase, or acetylcholinesterase; examples of suitable prosthetic group complexes include streptavidin/biotin and avidin/biotin; examples of suitable fluorescent materials include umbelliferone, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine fluorescein, dansyl chloride or phycoerythrin; an example of a luminescent material includes luminol; examples of bioluminescent materials include luciferase, luciferin, and aequorin; and examples of suitable radioactive material include iodine (121 I, 123 I, 125 I, 131 I), carbon (14 C), sulfur (35 S), tritium (3 H), indium (111 In, 112 In, 113 In, 115 In, 115 In), technetium (99 Tc, 99 Tc), thallium (201 Ti), gallium (68 Ga, 67 Ga), palladium (103 Pd), molybdenum (99 Mo), xenon (133 Xe), fluorine (18 F), 153 Sm, 177 Lu, 159 Gd, 149 Pm, 140 La, 175 Yb, 166 Ho, 90 Y, 47 Sc, 186 Re, 188 Re, 142 Pr, 105 Rh, and 97 Ru.

[0189] In specific embodiments, a polypeptide of the present invention or fragment or variant thereof is attached to macrocyclic chelators that associate with radiometal ions, including but not limited to, ¹⁷⁷Lu, ⁹⁰Y, ¹⁶⁶Ho, and ¹⁵³Sm, to polypeptides. In a preferred embodiment, the radiometal ion associated with the macrocyclic chelators is ¹¹¹In. In another preferred embodiment, the radiometal ion associated with the macrocyclic chelator is ⁹⁰Y. In specific embodiments, the macrocyclic chelator is 1,4,7,10-tetraazacyclododecane-N,N',N'',N'''-tetraacetic acid (DOTA). In other specific embodiments, DOTA is attached to an antibody of the invention or fragment thereof via a linker molecule. Examples of linker molecules useful for conjugating DOTA to a polypeptide are commonly known in the art - see, for example, DeNardo et al., Clin Cancer Res. 4(10):2483-90 (1998); Peterson et al., Bioconjug. Chem. 10(4):553-7 (1999); and Zimmerman et al, Nucl. Med. Biol. 26(8):943-50 (1999); which are hereby incorporated by reference in their entirety.

As mentioned, the immune/hematopoietic associated proteins of the invention [0190]may be modified by either natural processes, such as posttranslational processing, or by chemical modification techniques which are well known in the art. It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given immune/hematopoietic associated polypeptide. Immune/hematopoietic associated polypeptides may be branched, for example, as a result of ubiquitination, and they may be cyclic, with or without branching. Cyclic, branched, and branched cyclic immune/hematopoietic associated polypeptides may result from posttranslation natural processes or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphotidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cysteine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, pegylation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. (See, for instance, PROTEINS - STRUCTURE AND MOLECULAR PROPERTIES, 2nd

Ed., T. E. Creighton, W. H. Freeman and Company, New York (1993); POSTTRANSLATIONAL COVALENT MODIFICATION OF PROTEINS, B. C. Johnson, Ed., Academic Press, New York, pgs. 1-12 (1983); Seifter et al., Meth. Enzymol. 182:626-646 (1990); Rattan et al., Ann. N.Y. Acad. Sci. 663:48-62 (1992)).

- [0191] Also provided by the invention are chemically modified derivatives of the polypeptides of the invention which may provide additional advantages such as increased solubility, stability and circulating time of the polypeptide, or decreased immunogenicity (see U.S. Patent No. 4,179,337). The chemical moieties for derivitization may be selected from water soluble polymers such as polyethylene glycol, ethylene glycol/propylene glycol copolymers, carboxymethylcellulose, dextran, polyvinyl alcohol and the like. The polypeptides may be modified at random positions within the molecule, or at predetermined positions within the molecule and may include one, two, three or more attached chemical moieties.
- In polymer may be of any molecular weight, and may be branched or unbranched. For polyethylene glycol, the preferred molecular weight is between about 1 kDa and about 100 kDa (the term "about" indicating that in preparations of polyethylene glycol, some molecules will weigh more, some less, than the stated molecular weight) for ease in handling and manufacturing. Other sizes may be used, depending on the desired therapeutic profile (e.g., the duration of sustained release desired, the effects, if any on biological activity, the ease in handling, the degree or lack of antigenicity and other known effects of the polyethylene glycol to a therapeutic protein or analog). For example, the polyethylene glycol may have an average molecular weight of about 200, 500, 1000, 1500, 2000, 2500, 3000, 3500, 4000, 4500, 5000, 5500, 6000, 6500, 7000, 7500, 8000, 8500, 9000, 9500, 10,000, 10,500, 11,000, 11,500, 12,000, 12,500, 13,000, 13,500, 14,000, 14,500, 15,000, 15,500, 16,000, 16,500, 17,000, 17,500, 18,000, 18,500, 19,000, 19,500, 20,000, 25,000, 30,000, 35,000, 40,000, 50,000, 55,000, 60,000, 65,000, 70,000, 75,000, 80,000, 85,000, 90,000, 95,000, or 100,000 kDa.
- [0193] As noted above, the polyethylene glycol may have a branched structure. Branched polyethylene glycols are described, for example, in U.S. Patent No. 5,643,575; Morpurgo et al., Appl. Biochem. Biotechnol. 56:59-72 (1996); Vorobjev et al., Nucleosides Nucleotides 18:2745-2750 (1999); and Caliceti et al., Bioconjug. Chem. 10:638-646 (1999), the disclosures of each of which are incorporated herein by reference.

[0194] . The polyethylene glycol molecules (or other chemical moieties) should be attached to the protein with consideration of effects on functional or antigenic domains of the protein. There are a number of attachment methods available to those skilled in the art, such as, for example, the method disclosed in EP 0 401 384 (coupling PEG to G-CSF), herein incorporated by reference; see also Malik et al., Exp. Hematol. 20:1028-1035 (1992), reporting pegylation of GM-CSF using tresyl chloride. For example, polyethylene glycol may be covalently bound through amino acid residues via a reactive group, such as a free amino or carboxyl group. Reactive groups are those to which an activated polyethylene glycol molecule may be bound. The amino acid residues having a free amino group may include lysine residues and the N-terminal amino acid residues; those having a free carboxyl group may include aspartic acid residues glutamic acid residues and the C-terminal amino acid residue. Sulfhydryl groups may also be used as a reactive group for attaching the polyethylene glycol molecules. Preferred for therapeutic purposes is attachment at an amino group, such as attachment at the N-terminus or lysine group.

[0195] As suggested above, polyethylene glycol may be attached to proteins via linkage to any of a number of amino acid residues. For example, polyethylene glycol can be linked to proteins via covalent bonds to lysine, histidine, aspartic acid, glutamic acid, or cysteine residues. One or more reaction chemistries may be employed to attach polyethylene glycol to specific amino acid residues (e.g., lysine, histidine, aspartic acid, glutamic acid, or cysteine) of the protein or to more than one type of amino acid residue (e.g., lysine, histidine, aspartic acid, glutamic acid, cysteine and combinations thereof) of the protein.

Using polyethylene glycol as an illustration of the present composition, one may select from a variety of polyethylene glycol molecules (by molecular weight, branching, etc.), the proportion of polyethylene glycol molecules to protein (polypeptide) molecules in the reaction mix, the type of pegylation reaction to be performed, and the method of obtaining the selected N-terminally pegylated protein. The method of obtaining the N-terminally pegylated preparation (i.e., separating this moiety from other monopegylated moieties if necessary) may be by purification of the N-terminally pegylated material from a population of pegylated protein molecules. Selective proteins chemically modified at the N-terminus modification may be accomplished by reductive alkylation which exploits differential reactivity of different types of primary amino groups (lysine versus the

N-terminal) available for derivatization in a particular protein. Under the appropriate reaction conditions, substantially selective derivatization of the protein at the N-terminus with a carbonyl group containing polymer is achieved.

[0197] As indicated above, pegylation of the proteins of the invention may be accomplished by any number of means. For example, polyethylene glycol may be attached to the protein either directly or by an intervening linker. Linkerless systems for attaching polyethylene glycol to proteins are described in Delgado et al., Crit. Rev. Thera. Drug Carrier Sys. 9:249-304 (1992); Francis et al., Intern. J. of Hematol. 68:1-18 (1998); U.S. Patent No. 4,002,531; U.S. Patent No. 5,349,052; WO 95/06058; and WO 98/32466, the disclosures of each of which are incorporated herein by reference.

One system for attaching polyethylene glycol directly to amino acid residues of proteins without an intervening linker employs tresylated MPEG, which is produced by the modification of monmethoxy polyethylene glycol (MPEG) using tresylchloride (ClSO₂CH₂CF₃). Upon reaction of protein with tresylated MPEG, polyethylene glycol is directly attached to amine groups of the protein. Thus, the invention includes protein-polyethylene glycol conjugates produced by reacting proteins of the invention with a polyethylene glycol molecule having a 2,2,2-trifluoreothane sulphonyl group.

[0199] Polyethylene glycol can also be attached to proteins using a number of different intervening linkers. For example, U.S. Patent No. 5,612,460, the entire disclosure of which is incorporated herein by reference, discloses urethane linkers for connecting polyethylene glycol to proteins. Protein-polyethylene glycol conjugates wherein the polyethylene glycol is attached to the protein by a linker can also be produced by reaction of proteins with compounds such as MPEG-succinimidylsuccinate, MPEG activated with 1,1'-carbonyldiimidazole, MPEG-2,4,5-trichloropenylcarbonate, MPEG-p-nitrophenolcarbonate, and various MPEG-succinate derivatives. A number of additional polyethylene glycol derivatives and reaction chemistries for attaching polyethylene glycol to proteins are described in International Publication No. WO 98/32466, the entire disclosure of which is incorporated herein by reference. Pegylated protein products produced using the reaction chemistries set out herein are included within the scope of the invention.

[0200] The number of polyethylene glycol moieties attached to each protein of the invention (i.e., the degree of substitution) may also vary. For example, the pegylated

proteins of the invention may be linked, on average, to 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 12, 15, 17, 20, or more polyethylene glycol molecules. Similarly, the average degree of substitution within ranges such as 1-3, 2-4, 3-5, 4-6, 5-7, 6-8, 7-9, 8-10, 9-11, 10-12, 11-13, 12-14, 13-15, 14-16, 15-17, 16-18, 17-19, or 18-20 polyethylene glycol moieties per protein molecule. Methods for determining the degree of substitution are discussed, for example, in Delgado et al., Crit. Rev. Thera. Drug Carrier Sys. 9:249-304 (1992).

- [0201] The immune/hematopoietic associated polypeptides of the invention can be recovered and purified from chemical synthesis and recombinant cell cultures by standard methods which include, but are not limited to, ammonium sulfate or ethanol precipitation, acid extraction, anion or cation exchange chromatography, phosphocellulose chromatography, hydrophobic interaction chromatography, affinity chromatography, hydroxylapatite chromatography and lectin chromatography. Most preferably, high performance liquid chromatography ("HPLC") is employed for purification. Well known techniques for refolding protein may be employed to regenerate active conformation when the polypeptide is denatured during isolation and/or purification.
- [0202] Immune/hematopoietic associated polynucleotides and polypeptides may be used in accordance with the present invention for a variety of applications, particularly those that make use of the chemical and biological properties of immune/hematopoietic associated antigens. Among these are applications in the detection, prevention, diagnosis and/or treatment of diseases associated with hematopoiesis and/or the immune system, such as e.g., cancers and tumors of hematopoietic cells as well as diseases and disorders of/associated with hematopoiesis or the immune system (such as anemias, autoimmune diseases, immunodeficiencies, allergies, asthma, and inflammation) and infectious/parasitic diseases. Additional applications relate to diagnosis and to treatment of disorders of cells, tissues and organisms. These aspects of the invention are discussed further below.
- [0203] In a preferred embodiment, polynucleotides expressed in a particular tissue type (see, e.g., Table 1A, column 7) are used to detect, diagnose, treat, prevent and/or prognose disorders associated with the tissue type.
- [0204] The polypeptides of the invention may be in monomers or multimers (i.e., dimers, trimers, tetramers and higher multimers). Accordingly, the present invention relates to monomers and multimers of the polypeptides of the invention, their preparation,

and compositions (preferably, Therapeutics) containing them. In specific embodiments, the polypeptides of the invention are monomers, dimers, trimers or tetramers. In additional embodiments, the multimers of the invention are at least dimers, at least trimers, or at least tetramers.

[0205] Multimers encompassed by the invention may be homomers or heteromers. As used herein, the term homomer refers to a multimer containing only polypeptides corresponding to a protein of the invention (e.g., the amino acid sequence of SEQ ID NO:Y, an amino acid sequence encoded by SEQ ID NO:X or the complement of SEQ ID NO:X, the amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, and/or an amino acid sequence encoded by cDNA contained in Clone ID NO:Z (including fragments, variants, splice variants, and fusion proteins, corresponding to these as described herein)). These homomers may contain polypeptides having identical or different amino acid sequences. In a specific embodiment, a homomer of the invention is a multimer containing only polypeptides having an identical amino acid sequence. In another specific embodiment, a homomer of the invention is a multimer containing polypeptides having different amino acid sequences. In specific embodiments, the multimer of the invention is a homodimer (e.g., containing two polypeptides having identical or different amino acid sequences) or a homotrimer (e.g., containing three polypeptides having identical and/or different amino acid sequences). In additional embodiments, the homomeric multimer of the invention is at least a homodimer, at least a homotrimer, or at least a homotetramer.

[0206] As used herein, the term heteromer refers to a multimer containing two or more heterologous polypeptides (i.e., polypeptides of different proteins) in addition to the polypeptides of the invention. In a specific embodiment, the multimer of the invention is a heterodimer, a heterotrimer, or a heterotetramer. In additional embodiments, the heteromeric multimer of the invention is at least a heterodimer, at least a heterotrimer, or at least a heterotetramer.

[0207] Multimers of the invention may be the result of hydrophobic, hydrophilic, ionic and/or covalent associations and/or may be indirectly linked by, for example, liposome formation. Thus, in one embodiment, multimers of the invention, such as, for example, homodimers or homotrimers, are formed when polypeptides of the invention contact one another in solution. In another embodiment, heteromultimers of the invention, such as, for

example, heterotrimers or heterotetramers, are formed when polypeptides of the invention contact antibodies to the polypeptides of the invention (including antibodies to the heterologous polypeptide sequence in a fusion protein of the invention) in solution. In other embodiments, multimers of the invention are formed by covalent associations with and/or between the polypeptides of the invention. Such covalent associations may involve one or more amino acid residues contained in the polypeptide sequence (e.g., that recited in SEQ ID NO:Y, encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, and/or encoded by the cDNA contained in Clone ID NO:Z). In one instance, the covalent associations are cross-linking between cysteine residues located within the polypeptide sequences which interact in the native (i.e., naturally occurring) polypeptide. In another instance, the covalent associations are the consequence of chemical or recombinant manipulation. Alternatively, such covalent associations may involve one or more amino acid residues contained in the heterologous polypeptide sequence in a fusion protein. In one example, covalent associations are between the heterologous sequence contained in a fusion protein of the invention (see, e.g., U.S. Patent Number 5,478,925). In a specific example, the covalent associations are between the heterologous sequence contained in a Fc fusion protein of the invention (as described herein). In another specific example, covalent associations of fusion proteins of the invention are between heterologous polypeptide sequence from another protein that is capable of forming covalently associated multimers, such as for example, osteoprotegerin (see, e.g., International Publication NO: WO 98/49305, the contents of which are herein incorporated by reference in its entirety). In another embodiment, two or more polypeptides of the invention are joined through peptide linkers. Examples include those peptide linkers described in U.S. Pat. No. 5,073,627 (hereby incorporated by reference). Proteins comprising multiple polypeptides of the invention separated by peptide linkers may be produced using conventional recombinant DNA technology.

[0208] Another method for preparing multimer polypeptides of the invention involves use of polypeptides of the invention fused to a leucine zipper or isoleucine zipper polypeptide sequence. Leucine zipper and isoleucine zipper domains are polypeptides that promote multimerization of the proteins in which they are found. Leucine zippers were originally identified in several DNA-binding proteins (Landschulz et al., Science 240:1759, (1988)), and have since been found in a variety of different proteins. Among

the known leucine zippers are naturally occurring peptides and derivatives thereof that dimerize or trimerize. Examples of leucine zipper domains suitable for producing soluble multimeric proteins of the invention are those described in PCT application WO 94/10308, hereby incorporated by reference. Recombinant fusion proteins comprising a polypeptide of the invention fused to a polypeptide sequence that dimerizes or trimerizes in solution are expressed in suitable host cells, and the resulting soluble multimeric fusion protein is recovered from the culture supernatant using techniques known in the art.

[0209] Trimeric polypeptides of the invention may offer the advantage of enhanced biological activity. Preferred leucine zipper moieties and isoleucine moieties are those that preferentially form trimers. One example is a leucine zipper derived from lung surfactant protein D (SPD), as described in Hoppe et al. (FEBS Letters 344:191, (1994)) and in U.S. patent application Ser. No. 08/446,922, hereby incorporated by reference. Other peptides derived from naturally occurring trimeric proteins may be employed in preparing trimeric polypeptides of the invention.

[0210] In another example, proteins of the invention are associated by interactions between Flag® polypeptide sequence contained in fusion proteins of the invention containing Flag® polypeptide sequence. In a further embodiment, proteins of the invention are associated by interactions between heterologous polypeptide sequence contained in Flag® fusion proteins of the invention and anti-Flag® antibody.

known in the art. For example, polypeptides desired to be contained in the multimers of the invention may be chemically cross-linked using linker molecules and linker molecule length optimization techniques known in the art (see, e.g., U.S. Patent Number 5,478,925, which is herein incorporated by reference in its entirety). Additionally, multimers of the invention may be generated using techniques known in the art to form one or more intermolecule cross-links between the cysteine residues located within the sequence of the polypeptides desired to be contained in the multimer (see, e.g., U.S. Patent Number 5,478,925, which is herein incorporated by reference in its entirety). Further, polypeptides of the invention may be routinely modified by the addition of cysteine or biotin to the C-terminus or N-terminus of the polypeptide and techniques known in the art may be applied to generate multimers containing one or more of these modified polypeptides (see, e.g., U.S. Patent Number 5,478,925, which is herein incorporated by reference in its entirety).

Additionally, techniques known in the art may be applied to generate liposomes containing the polypeptide components desired to be contained in the multimer of the invention (see, e.g., U.S. Patent Number 5,478,925, which is herein incorporated by reference in its entirety).

[0212] Alternatively, multimers of the invention may be generated using genetic engineering techniques known in the art. In one embodiment, polypeptides contained in multimers of the invention are produced recombinantly using fusion protein technology described herein or otherwise known in the art (see, e.g., U.S. Patent Number 5,478,925, which is herein incorporated by reference in its entirety). In a specific embodiment, polynucleotides coding for a homodimer of the invention are generated by ligating a polynucleotide sequence encoding a polypeptide of the invention to a sequence encoding a linker polypeptide and then further to a synthetic polynucleotide encoding the translated product of the polypeptide in the reverse orientation from the original C-terminus to the N-terminus (lacking the leader sequence) (see, e.g., U..S Patent Number 5,478,925, which is herein incorporated by reference in its entirety). In another embodiment, recombinant techniques described herein or otherwise known in the art are applied to generate recombinant polypeptides of the invention which contain a transmembrane domain (or hydrophobic or signal peptide) and which can be incorporated by membrane reconstitution techniques into liposomes (see, e.g., U.S. Patent Number 5,478,925, which is herein incorporated by reference in its entirety).

Antibodies

Further polypeptides of the invention relate to antibodies and T-cell antigen receptors (TCR) which immunospecifically bind a polypeptide, polypeptide fragment, or variant of the invention (e.g., a polypeptide or fragment or variant of the amino acid sequence of SEQ ID NO:Y or a polypeptide encoded by the cDNA contained in Clone ID NO:Z, and/or an epitope, of the present invention) as determined by immunoassays well known in the art for assaying specific antibody-antigen binding. Antibodies of the invention include, but are not limited to, polyclonal, monoclonal, multispecific, human, humanized or chimeric antibodies, single chain antibodies, Fab fragments, F(ab') fragments, fragments produced by a Fab expression library, anti-idiotypic (anti-Id) antibodies (including, e.g., anti-Id antibodies to antibodies of the invention),

intracellularly-made antibodies (i.e., intrabodies), and epitope-binding fragments of any of the above. The term "antibody," as used herein, refers to immunoglobulin molecules and immunologically active portions of immunoglobulin molecules, i.e., molecules that contain an antigen binding site that immunospecifically binds an antigen. The immunoglobulin molecules of the invention can be of any type (e.g., IgG, IgE, IgM, IgD, IgA and IgY), class (e.g., IgG1, IgG2, IgG3, IgG4, IgA1 and IgA2) or subclass of immunoglobulin molecule. In preferred embodiments, the immunoglobulin molecules of the invention are IgG1. In other preferred embodiments, the immunoglobulin molecules of the invention are IgG4.

[0214]Most preferably the antibodies are human antigen-binding antibody fragments of the present invention and include, but are not limited to, Fab, Fab' and F(ab')2, Fd, single-chain Fvs (scFv), single-chain antibodies, disulfide-linked Fvs (sdFv) and fragments comprising either a VL or VH domain. Antigen-binding antibody fragments, including single-chain antibodies, may comprise the variable region(s) alone or in combination with the entirety or a portion of the following: hinge region, CH1, CH2, and CH3 domains. Also included in the invention are antigen-binding fragments also comprising any combination of variable region(s) with a hinge region, CH1, CH2, and CH3 domains. The antibodies of the invention may be from any animal origin including birds and mammals. Preferably, the antibodies are human, murine (e.g., mouse and rat). donkey, ship rabbit, goat, guinea pig, camel, horse, or chicken. As used herein, "human" antibodies include antibodies having the amino acid sequence of a human immunoglobulin and include antibodies isolated from human immunoglobulin libraries or from animals transgenic for one or more human immunoglobulin and that do not express endogenous immunoglobulins, as described infra and, for example in, U.S. Patent No. 5,939,598 by Kucherlapati et al.

[0215] The antibodies of the present invention may be monospecific, bispecific, trispecific or of greater multispecificity. Multispecific antibodies may be specific for different epitopes of a polypeptide of the present invention or may be specific for both a polypeptide of the present invention as well as for a heterologous epitope, such as a heterologous polypeptide or solid support material. See, e.g., PCT publications WO 93/17715; WO 92/08802; WO 91/00360; WO 92/05793; Tutt, et al., J. Immunol. 147:60-

69 (1991); U.S. Patent Nos. 4,474,893; 4,714,681; 4,925,648; 5,573,920; 5,601,819; Kostelny et al., J. Immunol. 148:1547-1553 (1992).

[0216] Antibodies of the present invention may be described or specified in terms of the epitope(s) or portion(s) of a polypeptide of the present invention which they recognize or specifically bind. The epitope(s) or polypeptide portion(s) may be specified as described herein, e.g., by N-terminal and C-terminal positions, or by size in contiguous amino acid residues, or listed in the Tables and Figures. Preferred epitopes of the invention include those shown in column 6 of Table 1A, as well as polynucleotides that encode these epitopes. Antibodies which specifically bind any epitope or polypeptide of the present invention may also be excluded. Therefore, the present invention includes antibodies that specifically bind polypeptides of the present invention, and allows for the exclusion of the same.

[0217]Antibodies of the present invention may also be described or specified in terms of their cross-reactivity. Antibodies that do not bind any other analog, ortholog, or homolog of a polypeptide of the present invention are included. Antibodies that bind polypeptides with at least 95%, at least 90%, at least 85%, at least 80%, at least 75%, at least 70%, at least 65%, at least 60%, at least 55%, and at least 50% identity (as calculated using methods known in the art and described herein) to a polypeptide of the present invention are also included in the present invention. In specific embodiments, antibodies of the present invention cross-react with murine, rat and/or rabbit homologs of human proteins and the corresponding epitopes thereof. Antibodies that do not bind polypeptides with less than 95%, less than 90%, less than 85%, less than 80%, less than 75%, less than 70%, less than 65%, less than 60%, less than 55%, and less than 50% identity (as calculated using methods known in the art and described herein) to a polypeptide of the present invention are also included in the present invention. In a specific embodiment, the above-described cross-reactivity is with respect to any single specific antigenic or immunogenic polypeptide, or combination(s) of 2, 3, 4, 5, or more of the specific antigenic and/or immunogenic polypeptides disclosed herein. Further included in the present invention are antibodies which bind polypeptides encoded by polynucleotides which hybridize to a polynucleotide of the present invention under stringent hybridization conditions (as described herein). Antibodies of the present invention may also be described or specified in terms of their binding affinity to a polypeptide of the invention.

Preferred binding affinities include those with a dissociation constant or Kd less than 5 X 10^{-2} M, 10^{-2} M, 5×10^{-3} M, 10^{-3} M, 5×10^{-4} M, 10^{-4} M, 5×10^{-5} M, 10^{-5} M, 5×10^{-6} M, 10^{-6} M, 5×10^{-7} M, 10^{7} M, 5×10^{-8} M, 10^{-8} M, 5×10^{-9} M, 10^{-9} M, 5×10^{-10} M, 10^{-10} M, 5×10^{-11} M, 10^{-11} M, 5×10^{-12} M, 10^{-12} M, 10^{-13} M, 10^{-13} M, 10^{-14} M, 10^{-14} M, 5×10^{-15} M, or 10^{-15} M.

[0218] The invention also provides antibodies that competitively inhibit binding of an antibody to an epitope of the invention as determined by any method known in the art for determining competitive binding, for example, the immunoassays described herei-n. In preferred embodiments, the antibody competitively inhibits binding to the epitope by at least 95%, at least 90%, at least 85 %, at least 80%, at least 75%, at least 70%, at least 60%, or at least 50%.

[0219]Antibodies of the present invention may act as agonists or antagonists of the polypeptides of the present invention. For example, the present invention includes antibodies which disrupt the receptor/ligand interactions with the polypeptides of the invention either partially or fully. Preferably, antibodies of the present invention bind an antigenic epitope disclosed herein, or a portion thereof. The invention features both receptor-specific antibodies and ligand-specific antibodies. The invention also features receptor-specific antibodies which do not prevent ligand binding but prevent receptor activation. Receptor activation (i.e., signaling) may be determined by techniques described herein or otherwise known in the art. For example, receptor activation can be determined by detecting the phosphorylation (e.g., tyrosine or serine/threonine) of the receptor or its substrate by immunoprecipitation followed by western blot analysis (for example, as described supra). In specific embodiments, antibodies are provided that inhibit ligand activity or receptor activity by at least 95%, at least 90%, at least 85%, at least 80%, at least 75%, at least 70%, at least 60%, or at least 50% of the activity in absence of the antibody.

[0220] The invention also features receptor-specific antibodies which both prevent ligand binding and receptor activation as well as antibodies that recognize the receptor-ligand complex, and, preferably, do not specifically recognize the unbound receptor or the unbound ligand. Likewise, included in the invention are neutralizing antibodies which bind the ligand and prevent binding of the ligand to the receptor, as well as antibodies which bind the ligand, thereby preventing receptor activation, but do not prevent the

ligand from binding the receptor. Further included in the invention are antibodies which activate the receptor. These antibodies may act as receptor agonists, i.e., potentiate or activate either all or a subset of the biological activities of the ligand-mediated receptor activation, for example, by inducing dimerization of the receptor. The antibodies may be specified as agonists, antagonists or inverse agonists for biological activities comprising the specific biological activities of the peptides of the invention disclosed herein. The above antibody agonists can be made using methods known in the art. See, e.g., PCT publication WO 96/40281; U.S. Patent No. 5,811,097; Deng et al., Blood 92(6):1981-1988 (1998); Chen et al., Cancer Res. 58(16):3668-3678 (1998); Harrop et al., J. Immunol. 161(4):1786-1794 (1998); Zhu et al., Cancer Res. 58(15):3209-3214 (1998); Yoon et al., J. Immunol. 160(7):3170-3179 (1998); Prat et al., J. Cell. Sci. 111(Pt2):237-247 (1998); Pitard et al., J. Immunol. Methods 205(2):177-190 (1997); Liautard et al., Cytokine 9(4):233-241 (1997); Carlson et al., J. Biol. Chem. 272(17):11295-11301 (1997); Taryman et al., Neuron 14(4):755-762 (1995); Muller et al., Structure 6(9):1153-1167 (1998); Bartunek et al., Cytokine 8(1):14-20 (1996) (which are all incorporated by reference herein in their entireties).

[0221] Antibodies of the present invention may be used, for example, to purify, detect, and target the polypeptides of the present invention, including both *in vitro* and *in vivo* diagnostic and therapeutic methods. For example, the antibodies have utility in immunoassays for qualitatively and quantitatively measuring levels of the polypeptides of the present invention in biological samples. See, e.g., Harlow et al., Antibodies: A Laboratory Manual, (Cold Spring Harbor Laboratory Press, 2nd ed. 1988); incorporated by reference herein in its entirety.

[0222] As discussed in more detail below, the antibodies of the present invention may be used either alone or in combination with other compositions. The antibodies may further be recombinantly fused to a heterologous polypeptide at the N- or C-terminus or chemically conjugated (including covalent and non-covalent conjugations) to polypeptides or other compositions. For example, antibodies of the present invention may be recombinantly fused or conjugated to molecules useful as labels in detection assays and effector molecules such as heterologous polypeptides, drugs, radionuclides, or toxins. See, e.g., PCT publications WO 92/08495; WO 91/14438; WO 89/12624; U.S. Patent No.

5,314,995; and EP 396,387; the disclosures of which are incorporated herein by reference in their entireties.

[0223] The antibodies of the invention include derivatives that are modified, i.e., by the covalent attachment of any type of molecule to the antibody such that covalent attachment does not prevent the antibody from generating an anti-idiotypic response. For example, but not by way of limitation, the antibody derivatives include antibodies that have been modified, e.g., by glycosylation, acetylation, pegylation, phosphylation, amidation, derivatization by known protecting/blocking groups, proteolytic cleavage, linkage to a cellular ligand or other protein, etc. Any of numerous chemical modifications may be carried out by known techniques, including, but not limited to specific chemical cleavage, acetylation, formylation, metabolic synthesis of tunicamycin, etc. Additionally, the derivative may contain one or more non-classical amino acids.

The antibodies of the present invention may be generated by any suitable method known in the art. Polyclonal antibodies to an antigen-of- interest can be produced by various procedures well known in the art. For example, a polypeptide of the invention can be administered to various host animals including, but not limited to, rabbits, mice, rats, etc. to induce the production of sera containing polyclonal antibodies specific for the antigen. Various adjuvants may be used to increase the immunological response, depending on the host species, and include but are not limited to, Freund's (complete and incomplete), mineral gels such as aluminum hydroxide, surface active substances such as lysolecithin, pluronic polyols, polyanions, peptides, oil emulsions, keyhole limpet hemocyanins, dinitrophenol, and potentially useful human adjuvants such as BCG (bacille Calmette-Guerin) and corynebacterium parvum. Such adjuvants are also well known in the art.

[0225] Monoclonal antibodies can be prepared using a wide variety of techniques known in the art including the use of hybridoma, recombinant, and phage display technologies, or a combination thereof. For example, monoclonal antibodies can be produced using hybridoma techniques including those known in the art and taught, for example, in Harlow et al., Antibodies: A Laboratory Manual, (Cold Spring Harbor Laboratory Press, 2nd ed. 1988); Hammerling, et al., in: Monoclonal Antibodies and T-Cell Hybridomas 563-681 (Elsevier, N.Y., 1981) (said references incorporated by reference in their entireties). The term "monoclonal antibody" as used herein is not

limited to antibodies produced through hybridoma technology. The term "monoclonal antibody" refers to an antibody that is derived from a single clone, including any eukaryotic, prokaryotic, or phage clone, and not the method by which it is produced.

Methods for producing and screening for specific antibodies using hybridoma technology are routine and well known in the art and are discussed in detail in the Examples. In a non-limiting example, mice can be immunized with a polypeptide of the invention or a cell expressing such peptide. Once an immune response is detected, e.g., antibodies specific for the antigen are detected in the mouse serum, the mouse spleen is harvested and splenocytes isolated. The splenocytes are then fused by well known techniques to any suitable myeloma cells, for example cells from cell line SP20 available from the ATCC. Hybridomas are selected and cloned by limited dilution. The hybridoma clones are then assayed by methods known in the art for cells that secrete antibodies capable of binding a polypeptide of the invention. Ascites fluid, which generally contains high levels of antibodies, can be generated by immunizing mice with positive hybridoma clones.

[0227] Accordingly, the present invention provides methods of generating monoclonal antibodies as well as antibodies produced by the method comprising culturing a hybridoma cell secreting an antibody of the invention wherein, preferably, the hybridoma is generated by fusing splenocytes isolated from a mouse immunized with an antigen of the invention with myeloma cells and then screening the hybridomas resulting from the fusion for hybridoma clones that secrete an antibody able to bind a polypeptide of the invention.

human B cell lines is transformation using Epstein Barr Virus (EBV). Protocols for generating EBV-transformed B cell lines are commonly known in the art, such as, for example, the protocol outlined in Chapter 7.22 of Current Protocols in Immunology, Coligan et al., Eds., 1994, John Wiley & Sons, NY, which is hereby incorporated in its entirety by reference herein. The source of B cells for transformation is commonly human peripheral blood, but B cells for transformation may also be derived from other sources including, but not limited to, lymph nodes, tonsil, spleen, tumor tissue, and infected tissues. Tissues are generally made into single cell suspensions prior to EBV transformation. Additionally, steps may be taken to either physically remove or inactivate

T cells (e.g., by treatment with cyclosporin A) in B cell-containing samples, because T cells from individuals seropositive for anti-EBV antibodies can suppress B cell immortalization by EBV.

[0229] In general, the sample containing human B cells is innoculated with EBV, and cultured for 3-4 weeks. A typical source of EBV is the culture supernatant of the B95-8 cell line (ATCC #VR-1492). Physical signs of EBV transformation can generally be seen towards the end of the 3-4 week culture period. By phase-contrast microscopy, transformed cells may appear large, clear, hairy and tend to aggregate in tight clusters of cells. Initially, EBV lines are generally polyclonal. However, over prolonged periods of cell cultures, EBV lines may become monoclonal or polyclonal as a result of the selective outgrowth of particular B cell clones. Alternatively, polyclonal EBV transformed lines may be subcloned (e.g., by limiting dilution culture) or fused with a suitable fusion partner and plated at limiting dilution to obtain monoclonal B cell lines. Suitable fusion partners for EBV transformed cell lines include mouse myeloma cell lines (e.g., SP2/0, X63-Ag8.653), heteromyeloma cell lines (human x mouse; e.g, SPAM-8, SBC-H20, and CB-F7), and human cell lines (e.g., GM 1500, SKO-007, RPMI 8226, and KR-4). Thus, the present invention also provides a method of generating polyclonal or monoclonal human antibodies against polypeptides of the invention or fragments thereof, comprising EBVtransformation of human B cells.

known techniques. For example, Fab and F(ab')2 fragments of the invention may be produced by proteolytic cleavage of immunoglobulin molecules, using enzymes such as papain (to produce Fab fragments) or pepsin (to produce F(ab')2 fragments). F(ab')2 fragments contain the variable region, the light chain constant region and the CH1 domain of the heavy chain. For example, the antibodies of the present invention can also be generated using various phage display methods known in the art and as discussed in detail in the Examples (e.g., Example 10). In phage display methods, functional antibody domains are displayed on the surface of phage particles which carry the polynucleotide sequences encoding them. In a particular embodiment, such phage can be utilized to display antigen binding domains expressed from a repertoire or combinatorial antibody library (e.g., human or murine). Phage expressing an antigen binding domain that binds the antigen of interest can be selected or identified with antigen, e.g., using labeled antigen

or antigen bound or captured to a solid surface or bead. Phage used in these methods are typically filamentous phage including fd and M13 binding domains expressed from phage with Fab, Fv or disulfide stabilized Fv antibody domains recombinantly fused to either the phage gene III or gene VIII protein. Examples of phage display methods that can be used to make the antibodies of the present invention include those disclosed in Brinkman et al., J. Immunol. Methods 182:41-50 (1995); Ames et al., J. Immunol. Methods 184:177-186 (1995); Kettleborough et al., Eur. J. Immunol. 24:952-958 (1994); Persic et al., Gene 187 9-18 (1997); Burton et al., Advances in Immunology 57:191-280 (1994); PCT application No. PCT/GB91/01134; PCT publications WO 90/02809; WO 91/10737; WO 92/01047; WO 92/18619; WO 93/11236; WO 95/15982; WO 95/20401; and U.S. Patent Nos. 5,698,426; 5,223,409; 5,403,484; 5,580,717; 5,427,908; 5,750,753; 5,821,047; 5,571,698; 5,427,908; 5,516,637; 5,780,225; 5,658,727; 5,733,743 and 5,969,108; each of which is incorporated herein by reference in its entirety.

[0231] As described in the above references, after phage selection, the antibody coding regions from the phage can be isolated and used to generate whole antibodies, including human antibodies, or any other desired antigen binding fragment, and expressed in any desired host, including mammalian cells, insect cells, plant cells, yeast, and bacteria, e.g., as described in detail below. For example, techniques to recombinantly produce Fab, Fab' and F(ab')2 fragments can also be employed using methods known in the art such as those disclosed in PCT publication WO 92/22324; Mullinax et al., BioTechniques 12(6):864-869 (1992); and Sawai et al., AJRI 34:26-34 (1995); and Better et al., Science 240:1041-1043 (1988) (said references incorporated by reference in their entireties).

[0232] Examples of techniques which can be used to produce single-chain Fvs and antibodies include those described in U.S. Patents 4,946,778 and 5,258,498; Huston et al., Methods in Enzymology 203:46-88 (1991); Shu et al., PNAS 90:7995-7999 (1993); and Skerra et al., Science 240:1038-1040 (1988). For some uses, including *in vivo* use of antibodies in humans and *in vitro* detection assays, it may be preferable to use chimeric, humanized, or human antibodies. A chimeric antibody is a molecule in which different portions of the antibody are derived from different animal species, such as antibodies having a variable region derived from a murine monoclonal antibody and a human immunoglobulin constant region. Methods for producing chimeric antibodies are known

in the art. See e.g., Morrison, Science 229:1202 (1985); Oi et al., BioTechniques 4:214 (1986); Gillies et al., (1989) J. Immunol. Methods 125:191-202; U.S. Patent Nos. 5,807,715; 4,816,567; and 4,816397, which are incorporated herein by reference in their Humanized antibodies are antibody molecules from non-human species antibody that binds the desired antigen having one or more complementarity determining regions (CDRs) from the non-human species and a framework regions from a human immunoglobulin molecule. Often, framework residues in the human framework regions will be substituted with the corresponding residue from the CDR donor antibody to alter, preferably improve, antigen binding. These framework substitutions are identified by methods well known in the art, e.g., by modeling of the interactions of the CDR and framework residues to identify framework residues important for antigen binding and sequence comparison to identify unusual framework residues at particular positions. (See, e.g., Queen et al., U.S. Patent No. 5,585,089; Riechmann et al., Nature 332:323 (1988), which are incorporated herein by reference in their entireties.) Antibodies can be humanized using a variety of techniques known in the art including, for example, CDRgrafting (EP 239,400; PCT publication WO 91/09967; U.S. Patent Nos. 5,225,539; 5,530,101; and 5,585,089), veneering or resurfacing (EP 592,106; EP 519,596; Padlan, Molecular Immunology 28(4/5):489-498 (1991); Studnicka et al., Protein Engineering 7(6):805-814 (1994); Roguska. et al., PNAS 91:969-973 (1994)), and chain shuffling (U.S. Patent No. 5,565,332).

- [0233] Completely human antibodies are particularly desirable for therapeutic treatment of human patients. Human antibodies can be made by a variety of methods known in the art including phage display methods described above using antibody libraries derived from human immunoglobulin sequences. See also, U.S. Patent Nos. 4,444,887 and 4,716,111; and PCT publications WO 98/46645, WO 98/50433, WO 98/24893, WO 98/16654, WO 96/34096, WO 96/33735, and WO 91/10741; each of which is incorporated herein by reference in its entirety.
- [0234] Human antibodies can also be produced using transgenic mice which are incapable of expressing functional endogenous immunoglobulins, but which can express human immunoglobulin genes. For example, the human heavy and light chain immunoglobulin gene complexes may be introduced randomly or by homologous recombination into mouse embryonic stem cells. Alternatively, the human variable

region, constant region, and diversity region may be introduced into mouse embryonic stem cells in addition to the human heavy and light chain genes. The mouse heavy and light chain immunoglobulin genes may be rendered non-functional separately or simultaneously with the introduction of human immunoglobulin loci by homologous recombination. In particular, homozygous deletion of the JH region prevents endogenous antibody production. The modified embryonic stem cells are expanded and microinjected into blastocysts to produce chimeric mice. The chimeric mice are then bred to produce homozygous offspring which express human antibodies. The transgenic mice are immunized in the normal fashion with a selected antigen, e.g., all or a portion of a polypeptide of the invention. Monoclonal antibodies directed against the antigen can be obtained from the immunized, transgenic mice using conventional hybridoma technology. The human immunoglobulin transgenes harbored by the transgenic mice rearrange during B cell differentiation, and subsequently undergo class switching and somatic mutation. Thus, using such a technique, it is possible to produce therapeutically useful IgG, IgA, IgM and IgE antibodies. For an overview of this technology for producing human antibodies, see Lonberg and Huszar, Int. Rev. Immunol. 13:65-93 (1995). For a detailed discussion of this technology for producing human antibodies and human monoclonal antibodies and protocols for producing such antibodies, see, e.g., PCT publications WO 98/24893; WO 92/01047; WO 96/34096; WO 96/33735; European Patent No. 0 598 877; U.S. Patent Nos. 5,413,923; 5,625,126; 5,633,425; 5,569,825; 5,661,016; 5,545,806; 5,814,318; 5,885,793; 5,916,771; 5,939,598; 6,075,181 and 6,114,598, which are incorporated by reference herein in their entirety. In addition, companies such as Abgenix, Inc. (Freemont, CA) and Genpharm (San Jose, CA) can be engaged to provide human antibodies directed against a selected antigen using technology similar to that described above.

[0235] Completely human antibodies which recognize a selected epitope can be generated using a technique referred to as "guided selection." In this approach a selected non-human monoclonal antibody, e.g., a mouse antibody, is used to guide the selection of a completely human antibody recognizing the same epitope. (Jespers et al., Bio/technology 12:899-903 (1988)).

[0236] Further, antibodies to the polypeptides of the invention can, in turn, be utilized to generate anti-idiotype antibodies that "mimic" polypeptides of the invention using

techniques well known to those skilled in the art. (See, e.g., Greenspan & Bona, FASEB J. 7(5):437-444; (1989) and Nissinoff, J. Immunol. 147(8):2429-2438 (1991)). For example, antibodies which bind to and competitively inhibit polypeptide multimerization and/or binding of a polypeptide of the invention to a ligand can be used to generate antiidiotypes that "mimic" the polypeptide multimerization and/or binding domain and, as a consequence, bind to and neutralize polypeptide and/or its ligand. Such neutralizing antiidiotypes or Fab fragments of such anti-idiotypes can be used in therapeutic regimens to neutralize polypeptide ligand/receptor. For example, such anti-idiotypic antibodies can be used to bind a polypeptide of the invention and/or to bind its ligand(s)/receptor(s), and thereby block its biological activity. Alternatively, antibodies which bind to and enhance polypeptide multimerization and/or binding, and/or receptor/ligand multimerization, binding and/or signaling can be used to generate anti-idiotypes that function as agonists of a polypeptide of the invention and/or its ligand/receptor. Such agonistic anti-idiotypes or Fab fragments of such anti-idiotypes can be used in therapeutic regimens as agonists of the polypeptides of the invention or its ligand(s)/receptor(s). For example, such antiidiotypic antibodies can be used to bind a polypeptide of the invention and/or to bind its ligand(s)/receptor(s), and thereby promote or enhance its biological activity.

Intrabodies of the invention can be produced using methods known in the art, such as those disclosed and reviewed in Chen et al., Hum. Gene Ther. 5:595-601 (1994); Marasco, W.A., Gene Ther. 4:11-15 (1997); Rondon and Marasco, Annu. Rev. Microbiol. 51:257-283 (1997); Proba et al., J. Mol. Biol. 275:245-253 (1998); Cohen et al., Oncogene 17:2445-2456 (1998); Ohage and Steipe, J. Mol. Biol. 291:1119-1128 (1999); Ohage et al., J. Mol. Biol. 291:1129-1134 (1999); Wirtz and Steipe, Protein Sci. 8:2245-2250 (1999); Zhu et al., J. Immunol. Methods 231:207-222 (1999); and references cited therein.

Polynucleotides Encoding Antibodies

[0238] The invention further provides polynucleotides comprising a nucleotide sequence encoding an antibody of the invention and fragments thereof. The invention also encompasses polynucleotides that hybridize under stringent or alternatively, under lower stringency hybridization conditions, e.g., as defined *supra*, to polynucleotides that encode an antibody, preferably, that specifically binds to a polypeptide of the invention, preferably, an antibody that binds to a polypeptide having the amino acid sequence of

SEQ ID NO:Y, to a polypeptide encoded by a portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, and/or to a polypeptide encoded by the cDNA contained in Clone ID NO:Z.

[0239] The polynucleotides may be obtained, and the nucleotide sequence of the polynucleotides determined, by any method known in the art. For example, if the nucleotide sequence of the antibody is known, a polynucleotide encoding the antibody may be assembled from chemically synthesized oligonucleotides (e.g., as described in Kutmeier et al., BioTechniques 17:242 (1994)), which, briefly, involves the synthesis of overlapping oligonucleotides containing portions of the sequence encoding the antibody, annealing and ligating of those oligonucleotides, and then amplification of the ligated oligonucleotides by PCR.

[0240] Alternatively, a polynucleotide encoding an antibody may be generated from nucleic acid from a suitable source. If a clone containing a nucleic acid encoding a particular antibody is not available, but the sequence of the antibody molecule is known, a nucleic acid encoding the immunoglobulin may be chemically synthesized or obtained from a suitable source (e.g., an antibody cDNA library, or a cDNA library generated from, or nucleic acid, preferably poly A+RNA, isolated from, any tissue or cells expressing the antibody, such as hybridoma cells selected to express an antibody of the invention) by PCR amplification using synthetic primers hybridizable to the 3' and 5' ends of the sequence or by cloning using an oligonucleotide probe specific for the particular gene sequence to identify, e.g., a cDNA clone from a cDNA library that encodes the antibody. Amplified nucleic acids generated by PCR may then be cloned into replicable cloning vectors using any method well known in the art.

[0241] Once the nucleotide sequence and corresponding amino acid sequence of the antibody is determined, the nucleotide sequence of the antibody may be manipulated using methods well known in the art for the manipulation of nucleotide sequences, e.g., recombinant DNA techniques, site directed mutagenesis, PCR, etc. (see, for example, the techniques described in Sambrook et al., 1990, Molecular Cloning, A Laboratory Manual, 2d Ed., Cold Spring Harbor Laboratory, Cold Spring Harbor, NY and Ausubel et al., eds., 1998, Current Protocols in Molecular Biology, John Wiley & Sons, NY, which are both incorporated by reference herein in their entireties), to generate antibodies having a

different amino acid sequence, for example to create amino acid substitutions, deletions, and/or insertions.

In a specific embodiment, the amino acid sequence of the heavy and/or light [0242] chain variable domains may be inspected to identify the sequences of the complementarity determining regions (CDRs) by methods that are well know in the art, e.g., by comparison to known amino acid sequences of other heavy and light chain variable regions to determine the regions of sequence hypervariability. Using routine recombinant DNA techniques, one or more of the CDRs may be inserted within framework regions, e.g., into human framework regions to humanize a non-human antibody, as described supra. The framework regions may be naturally occurring or consensus framework regions, and preferably human framework regions (see, e.g., Chothia et al., J. Mol. Biol. 278: 457-479 (1998) for a listing of human framework regions). Preferably, the polynucleotide generated by the combination of the framework regions and CDRs encodes an antibody that specifically binds a polypeptide of the invention. Preferably, as discussed supra, one or more amino acid substitutions may be made within the framework regions, and, preferably, the amino acid substitutions improve binding of the antibody to its antigen. Additionally, such methods may be used to make amino acid substitutions or deletions of one or more variable region cysteine residues participating in an intrachain disulfide bond to generate antibody molecules lacking one or more intrachain disulfide bonds. Other alterations to the polynucleotide are encompassed by the present invention and within the skill of the art.

[0243] In addition, techniques developed for the production of "chimeric antibodies" (Morrison et al., Proc. Natl. Acad. Sci. 81:851-855 (1984); Neuberger et al., Nature 312:604-608 (1984); Takeda et al., Nature 314:452-454 (1985)) by splicing genes from a mouse antibody molecule of appropriate antigen specificity together with genes from a human antibody molecule of appropriate biological activity can be used. As described supra, a chimeric antibody is a molecule in which different portions are derived from different animal species, such as those having a variable region derived from a murine mAb and a human immunoglobulin constant region, e.g., humanized antibodies.

[0244] Alternatively, techniques described for the production of single chain antibodies (U.S. Patent No. 4,946,778; Bird, Science 242:423- 42 (1988); Huston et al., Proc. Natl. Acad. Sci. USA 85:5879-5883 (1988); and Ward et al., Nature 334:544-54

(1989)) can be adapted to produce single chain antibodies. Single chain antibodies are formed by linking the heavy and light chain fragments of the Fv region via an amino acid bridge, resulting in a single chain polypeptide. Techniques for the assembly of functional Fv fragments in *E. coli* may also be used (Skerra et al., Science 242:1038-1041 (1988)).

Methods of Producing Antibodies

[0245] The antibodies of the invention can be produced by any method known in the art for the synthesis of antibodies, in particular, by chemical synthesis or preferably, by recombinant expression techniques. Methods of producing antibodies include, but are not limited to, hybridoma technology, EBV transformation, and other methods discussed herein as well as through the use recombinant DNA technology, as discussed below.

[0246] Recombinant expression of an antibody of the invention, or fragment, derivative or analog thereof, (e.g., a heavy or light chain of an antibody of the invention or a single chain antibody of the invention), requires construction of an expression vector containing a polynucleotide that encodes the antibody. Once a polynucleotide encoding an antibody molecule or a heavy or light chain of an antibody, or portion thereof (preferably containing the heavy or light chain variable domain), of the invention has been obtained, the vector for the production of the antibody molecule may be produced by recombinant DNA technology using techniques well known in the art. Thus, methods for preparing a protein by expressing a polynucleotide containing an antibody encoding nucleotide sequence are described herein. Methods which are well known to those skilled in the art can be used to construct expression vectors containing antibody coding sequences and appropriate transcriptional and translational control signals. These methods include, for example, in vitro recombinant DNA techniques, synthetic techniques, and in vivo genetic recombination. The invention, thus, provides replicable vectors comprising a nucleotide sequence encoding an antibody molecule of the invention, or a heavy or light chain thereof, or a heavy or light chain variable domain, operably linked to a promoter. Such vectors may include the nucleotide sequence encoding the constant region of the antibody molecule (see, e.g., PCT Publication WO 86/05807; PCT Publication WO 89/01036; and U.S. Patent No. 5,122,464) and the variable domain of the antibody may be cloned into such a vector for expression of the entire heavy or light chain.

and the transfected cells are then cultured by conventional techniques to produce an antibody of the invention. Thus, the invention includes host cells containing a polynucleotide encoding an antibody of the invention, or a heavy or light chain thereof, or a single chain antibody of the invention, operably linked to a heterologous promoter. In preferred embodiments for the expression of double-chained antibodies, vectors encoding both the heavy and light chains may be co-expressed in the host cell for expression of the entire immunoglobulin molecule, as detailed below.

A variety of host-expression vector systems may be utilized to express the [0248] antibody molecules of the invention. Such host-expression systems represent vehicles by which the coding sequences of interest may be produced and subsequently purified, but also represent cells which may, when transformed or transfected with the appropriate nucleotide coding sequences, express an antibody molecule of the invention in situ. These include but are not limited to microorganisms such as bacteria (e.g., E. coli, B. subtilis) transformed with recombinant bacteriophage DNA, plasmid DNA or cosmid DNA expression vectors containing antibody coding sequences; yeast (e.g., Saccharomyces, Pichia) transformed with recombinant yeast expression vectors containing antibody coding sequences; insect cell systems infected with recombinant virus expression vectors (e.g., baculovirus) containing antibody coding sequences; plant cell systems infected with recombinant virus expression vectors (e.g., cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or transformed with recombinant plasmid expression vectors (e.g., Ti plasmid) containing antibody coding sequences; or mammalian cell systems (e.g., COS, CHO. BHK. 293, 3T3 cells) harboring recombinant expression constructs containing promoters derived from the genome of mammalian cells (e.g., metallothionein promoter) or from mammalian viruses (e.g., the adenovirus late promoter; the vaccinia virus 7.5K promoter). Preferably, bacterial cells such as Escherichia coli, and more preferably, eukaryotic cells, especially for the expression of whole recombinant antibody molecule, are used for the expression of a recombinant antibody molecule. For example, mammalian cells such as Chinese hamster ovary cells (CHO), in conjunction with a vector such as the major intermediate early gene promoter element from human cytomegalovirus is an effective expression system for antibodies (Foecking et al., Gene 45:101 (1986); Cockett et al., Bio/Technology 8:2 (1990)).

[0249] In bacterial systems, a number of expression vectors may be advantageously selected depending upon the use intended for the antibody molecule being expressed. For example, when a large quantity of such a protein is to be produced, for the generation of pharmaceutical compositions of an antibody molecule, vectors which direct the expression of high levels of fusion protein products that are readily purified may be desirable. Such vectors include, but are not limited, to the E. coli expression vector pUR278 (Ruther et al., EMBO J. 2:1791 (1983)), in which the antibody coding sequence may be ligated individually into the vector in frame with the lac Z coding region so that a fusion protein is produced; pIN vectors (Inouye & Inouye, Nucleic Acids Res. 13:3101-3109 (1985); Van Heeke & Schuster, J. Biol. Chem. 24:5503-5509 (1989)); and the like. pGEX vectors may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such fusion proteins are soluble and can easily be purified from lysed cells by adsorption and binding to matrix glutathioneagarose beads followed by elution in the presence of free glutathione. The pGEX vectors are designed to include thrombin or factor Xa protease cleavage sites so that the cloned target gene product can be released from the GST moiety.

[0250] In an insect system, Autographa californica nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes. The virus grows in *Spodoptera frugiperda* cells. The antibody coding sequence may be cloned individually into non-essential regions (for example the polyhedrin gene) of the virus and placed under control of an AcNPV promoter (for example the polyhedrin promoter).

In mammalian host cells, a number of viral-based expression systems may be utilized. In cases where an adenovirus is used as an expression vector, the antibody coding sequence of interest may be ligated to an adenovirus transcription/translation control complex, e.g., the late promoter and tripartite leader sequence. This chimeric gene may then be inserted in the adenovirus genome by *in vitro* or *in vivo* recombination. Insertion in a non- essential region of the viral genome (e.g., region E1 or E3) will result in a recombinant virus that is viable and capable of expressing the antibody molecule in infected hosts. (e.g., see Logan & Shenk, Proc. Natl. Acad. Sci. USA 81:355-359 (1984)). Specific initiation signals may also be required for efficient translation of inserted antibody coding sequences. These signals include the ATG initiation codon and adjacent sequences. Furthermore, the initiation codon must be in phase with the reading frame of

translational control signals and initiation codons can be of a variety of origins, both natural and synthetic. The efficiency of expression may be enhanced by the inclusion of appropriate transcription enhancer elements, transcription terminators, etc. (see Bittner et al., Methods in Enzymol. 153:51-544 (1987)).

In addition, a host cell strain may be chosen which modulates the expression of the inserted sequences, or modifies and processes the gene product in the specific fashion desired. Such modifications (e.g., glycosylation) and processing (e.g., cleavage) of protein products may be important for the function of the protein. Different host cells have characteristic and specific mechanisms for the post-translational processing and modification of proteins and gene products. Appropriate cell lines or host systems can be chosen to ensure the correct modification and processing of the foreign protein expressed. To this end, eukaryotic host cells which possess the cellular machinery for proper processing of the primary transcript, glycosylation, and phosphorylation of the gene product may be used. Such mammalian host cells include but are not limited to CHO, VERY, BHK, Hela, COS, MDCK, 293, 3T3, WI38, and in particular, breast cancer cell lines such as, for example, BT483, Hs578T, HTB2, BT20 and T47D, and normal mammary gland cell line such as, for example, CRL7030 and Hs578Bst.

[0253] For long-term, high-yield production of recombinant proteins, stable expression is preferred. For example, cell lines which stably express the antibody molecule may be engineered. Rather than using expression vectors which contain viral origins of replication, host cells can be transformed with DNA controlled by appropriate expression control elements (e.g., promoter, enhancer, sequences, transcription terminators, polyadenylation sites, etc.), and a selectable marker. Following the introduction of the foreign DNA, engineered cells may be allowed to grow for 1-2 days in an enriched media, and then are switched to a selective media. The selectable marker in the recombinant plasmid confers resistance to the selection and allows cells to stably integrate the plasmid into their chromosomes and grow to form foci which in turn can be cloned and expanded into cell lines. This method may advantageously be used to engineer cell lines which express the antibody molecule. Such engineered cell lines may be particularly useful in screening and evaluation of compounds that interact directly or indirectly with the antibody molecule.

[0254] A number of selection systems may be used, including but not limited to the herpes simplex virus thymidine kinase (Wigler et al., Cell 11:223 (1977)), hypoxanthineguanine phosphoribosyltransferase (Szybalska & Szybalski, Proc. Natl. Acad. Sci. USA 48:202 (1992)), and adenine phosphoribosyltransferase (Lowy et al., Cell 22:817 (1980)) genes can be employed in tk-, hgprt- or aprt- cells, respectively. Also, antimetabolite resistance can be used as the basis of selection for the following genes: dhfr, which confers resistance to methotrexate (Wigler et al., Natl. Acad. Sci. USA 77:357 (1980); O'Hare et al., Proc. Natl. Acad. Sci. USA 78:1527 (1981)); gpt, which confers resistance to mycophenolic acid (Mulligan & Berg, Proc. Natl. Acad. Sci. USA 78:2072 (1981)); neo, which confers resistance to the aminoglycoside G-418 Clinical Pharmacy 12:488-505; Wu and Wu, Biotherapy 3:87-95 (1991); Tolstoshev, Ann. Rev. Pharmacol. Toxicol. 32:573-596 (1993); Mulligan, Science 260:926-932 (1993); and Morgan and Anderson, Ann. Rev. Biochem. 62:191-217 (1993); TIB TECH 11(5):155-215 (1993)); and hygro, which confers resistance to hygromycin (Santerre et al., Gene 30:147 (1984)). Methods commonly known in the art of recombinant DNA technology may be routinely applied to select the desired recombinant clone, and such methods are described, for example, in Ausubel et al. (eds.), Current Protocols in Molecular Biology, John Wiley & Sons, NY (1993); Kriegler, Gene Transfer and Expression, A Laboratory Manual, Stockton Press, NY (1990); and in Chapters 12 and 13, Dracopoli et al. (eds), Current Protocols in Human Genetics, John Wiley & Sons, NY (1994); Colberre-Garapin et al., J. Mol. Biol. 150:1 (1981), which are incorporated by reference herein in their entireties.

[0255] The expression levels of an antibody molecule can be increased by vector amplification (for a review, see Bebbington and Hentschel, The use of vectors based on gene amplification for the expression of cloned genes in mammalian cells in DNA cloning, Vol.3. (Academic Press, New York, 1987)). When a marker in the vector system expressing antibody is amplifiable, increase in the level of inhibitor present in culture of host cell will increase the number of copies of the marker gene. Since the amplified region is associated with the antibody gene, production of the antibody will also increase (Crouse et al., Mol. Cell. Biol. 3:257 (1983)).

[0256] Vectors which use glutamine synthase (GS) or DHFR as the selectable markers can be amplified in the presence of the drugs methionine sulphoximine or methotrexate, respectively. An advantage of glutamine synthase based vectors are the availabilty of cell

lines (e.g., the murine myeloma cell line, NS0) which are glutamine synthase negative. Glutamine synthase expression systems can also function in glutamine synthase expressing cells (e.g., Chinese Hamster Ovary (CHO) cells) by providing additional inhibitor to prevent the functioning of the endogenous gene. A glutamine synthase expression system and components thereof are detailed in PCT publications: WO87/04462; WO86/05807; WO89/01036; WO89/10404; and WO91/06657 which are incorporated in their entireties by reference herein. Additionally, glutamine synthase expression vectors that may be used according to the present invention are commercially available from suplliers, including, for example Lonza Biologics, Inc. (Portsmouth, NH). Expression and production of monoclonal antibodies using a GS expression system in murine myeloma cells is described in Bebbington et al., Bio/technology 10:169(1992) and in Biblia and Robinson Biotechnol. Prog. 11:1 (1995) which are incorporated in their entirities by reference herein.

[0257] The host cell may be co-transfected with two expression vectors of the invention, the first vector encoding a heavy chain derived polypeptide and the second vector encoding a light chain derived polypeptide. The two vectors may contain identical selectable markers which enable equal expression of heavy and light chain polypeptides. Alternatively, a single vector may be used which encodes, and is capable of expressing, both heavy and light chain polypeptides. In such situations, the light chain should be placed before the heavy chain to avoid an excess of toxic free heavy chain (Proudfoot, Nature 322:52 (1986); Kohler, Proc. Natl. Acad. Sci. USA 77:2197 (1980)). The coding sequences for the heavy and light chains may comprise cDNA or genomic DNA.

[0258] Once an antibody molecule of the invention has been produced by an animal, chemically synthesized, or recombinantly expressed, it may be purified by any method known in the art for purification of an immunoglobulin molecule, for example, by chromatography (e.g., ion exchange, affinity, particularly by affinity for the specific antigen after Protein A, and sizing column chromatography), centrifugation, differential solubility, or by any other standard technique for the purification of proteins. In addition, the antibodies of the present invention or fragments thereof can be fused to heterologous polypeptide sequences described herein or otherwise known in the art, to facilitate purification.

[0259] The present invention encompasses antibodies recombinantly fused or chemically conjugated (including both covalently and non-covalently conjugations) to a polypeptide (or portion thereof, preferably at least 10, 20, 30, 40, 50, 60, 70, 80, 90 or 100 amino acids of the polypeptide) of the present invention to generate fusion proteins. The fusion does not necessarily need to be direct, but may occur through linker sequences. The antibodies may be specific for antigens other than polypeptides (or portion thereof, preferably at least 10, 20, 30, 40, 50, 60, 70, 80, 90 or 100 amino acids of the polypeptide) of the present invention. For example, antibodies may be used to target the polypeptides of the present invention to particular cell types, either in vitro or in vivo, by fusing or conjugating the polypeptides of the present invention to antibodies specific for particular cell surface receptors. Antibodies fused or conjugated to the polypeptides of the present invention may also be used in in vitro immunoassays and purification methods using methods known in the art. See e.g., Harbor et al., supra, and PCT publication WO 93/21232; EP 439,095; Naramura et al., Immunol. Lett. 39:91-99 (1994); U.S. Patent 5,474,981; Gillies et al., PNAS 89:1428-1432 (1992); Fell et al., J. Immunol. 146:2446-2452 (1991), which are incorporated by reference in their entireties.

The present invention further includes compositions comprising the [0260] polypeptides of the present invention fused or conjugated to antibody domains other than the variable regions. For example, the polypeptides of the present invention may be fused or conjugated to an antibody Fc region, or portion thereof. The antibody portion fused to a polypeptide of the present invention may comprise the constant region, hinge region, CH1 domain, CH2 domain, and CH3 domain or any combination of whole domains or portions thereof. The polypeptides may also be fused or conjugated to the above antibody portions to form multimers. For example, Fc portions fused to the polypeptides of the present invention can form dimers through disulfide bonding between the Fc portions. Higher multimeric forms can be made by fusing the polypeptides to portions of IgA and IgM. Methods for fusing or conjugating the polypeptides of the present invention to antibody portions are known in the art. See, e.g., U.S. Patent Nos. 5,336,603; 5,622,929; 5,359,046; 5,349,053; 5,447,851; 5,112,946; EP 307,434; EP 367,166; PCT publications WO 96/04388; WO 91/06570; Ashkenazi et al., Proc. Natl. Acad. Sci. USA 88:10535-10539 (1991); Zheng et al., J. Immunol. 154:5590-5600 (1995); and Vil et al., Proc. Natl.

Acad. Sci. USA 89:11337- 11341 (1992) (said references incorporated by reference in their entireties).

As discussed, supra, the polypeptides corresponding to a polypeptide, [0261] polypeptide fragment, or a variant of SEQ ID NO:Y may be fused or conjugated to the above antibody portions to increase the in vivo half life of the polypeptides or for use in immunoassays using methods known in the art. Further, the polypeptides corresponding to SEQ ID NO:Y may be fused or conjugated to the above antibody portions to facilitate purification. One reported example describes chimeric proteins consisting of the first two domains of the human CD4-polypeptide and various domains of the constant regions of the heavy or light chains of mammalian immunoglobulins. See EP 394,827; Traunecker et al., Nature 331:84-86 (1988). The polypeptides of the present invention fused or conjugated to an antibody having disulfide- linked dimeric structures (due to the IgG) may also be more efficient in binding and neutralizing other molecules, than the monomeric secreted protein or protein fragment alone. See, for example, Fountoulakis et al., J. Biochem. 270:3958-3964 (1995). In many cases, the Fc part in a fusion protein is beneficial in therapy and diagnosis, and thus can result in, for example, improved pharmacokinetic properties. See, for example, EP A 232,262. Alternatively, deleting the Fc part after the fusion protein has been expressed, detected, and purified, would be desired. For example, the Fc portion may hinder therapy and diagnosis if the fusion protein is used as an antigen for immunizations. In drug discovery, for example, human proteins, such as hIL-5, have been fused with Fc portions for the purpose of highthroughput screening assays to identify antagonists of hIL-5. (See, Bennett et al., J. Molecular Recognition 8:52-58 (1995); Johanson et al., J. Biol. Chem. 270:9459-9471 (1995)).

[0262] Moreover, the antibodies or fragments thereof of the present invention can be fused to marker sequences, such as a peptide to facilitate purification. In preferred embodiments, the marker amino acid sequence is a hexa-histidine peptide, such as the tag provided in a pQE vector (QIAGEN, Inc., 9259 Eton Avenue, Chatsworth, CA, 91311), among others, many of which are commercially available. As described in Gentz et al., Proc. Natl. Acad. Sci. USA 86:821-824 (1989), for instance, hexa-histidine provides for convenient purification of the fusion protein. Other peptide tags useful for purification include, but are not limited to, the "HA" tag, which corresponds to an epitope derived

from the influenza hemagglutinin protein (Wilson et al., Cell 37:767 (1984)) and the "flag" tag.

[0263] The present invention further encompasses antibodies or fragments thereof conjugated to a diagnostic or therapeutic agent. The antibodies can be used diagnostically to, for example, monitor the development or progression of a tumor as part of a clinical testing procedure to, e.g., determine the efficacy of a given treatment regimen. Detection can be facilitated by coupling the antibody to a detectable substance. Examples of detectable substances include various enzymes, prosthetic groups, fluorescent materials, luminescent materials, bioluminescent materials, radioactive materials, positron emitting metals using various positron emission tomographies, and nonradioactive paramagnetic metal ions. The detectable substance may be coupled or conjugated either directly to the antibody (or fragment thereof) or indirectly, through an intermediate (such as, for example, a linker known in the art) using techniques known in the art. See, for example, U.S. Patent No. 4,741,900 for metal ions which can be conjugated to antibodies for use as diagnostics according to the present invention.

Further, an antibody or fragment thereof may be conjugated to a therapeutic [0264] moiety such as a cytotoxin, e.g., a cytostatic or cytocidal agent, a therapeutic agent or a radioactive metal ion, e.g., alpha-emitters such as, for example, 213Bi. A cytotoxin or cytotoxic agent includes any agent that is detrimental to cells. Examples include paclitaxol, cytochalasin B, gramicidin D, ethidium bromide, emetine, mitomycin, etoposide, tenoposide, vincristine, vinblastine, colchicin, doxorubicin, daunorubicin, dihydroxy anthracin dione, mitoxantrone, mithramycin, actinomycin D, 1dehydrotestosterone, glucocorticoids, procaine, tetracaine, lidocaine, propranolol, and puromycin and analogs or homologs thereof. Therapeutic agents include, but are not limited to, antimetabolites (e.g., methotrexate, 6-mercaptopurine, 6-thioguanine, cytarabine, 5-fluorouracil decarbazine), alkylating agents (e.g., mechlorethamine, thioepa chlorambucil, melphalan, carmustine (BSNU) and lomustine (CCNU), cyclothosphamide, busulfan, dibromomannitol, streptozotocin, mitomycin C, and cis- dichlorodiamine platinum (II) (DDP) cisplatin), anthracyclines (e.g., daunorubicin (formerly daunomycin) and doxorubicin), antibiotics (e.g., dactinomycin (formerly actinomycin), bleomycin, mithramycin, and anthramycin (AMC)), and anti-mitotic agents (e.g., vincristine and vinblastine).

The conjugates of the invention can be used for modifying a given biological response, the therapeutic agent or drug moiety is not to be construed as limited to classical chemical therapeutic agents. For example, the drug moiety may be a protein or polypeptide possessing a desired biological activity. Such proteins may include, for example, a toxin such as abrin, ricin A, pseudomonas exotoxin, or diphtheria toxin; a protein such as tumor necrosis factor, a-interferon, β-interferon, nerve growth factor, platelet derived growth factor, tissue plasminogen activator, an apoptotic agent, e.g., TNF-alpha, TNF-beta, AIM I (See, International Publication No. WO 97/33899), AIM II (See, International Publication No. WO 97/34911), Fas Ligand (Takahashi *et al.*, *Int. Immunol.*, 6:1567-1574 (1994)), VEGI (See, International Publication No. WO 99/23105), a thrombotic agent or an anti- angiogenic agent, e.g., angiostatin or endostatin; or, biological response modifiers such as, for example, lymphokines, interleukin-1 ("IL-1"), interleukin-2 ("IL-2"), interleukin-6 ("IL-6"), granulocyte macrophage colony stimulating factor ("GM-CSF"), granulocyte colony stimulating factor ("G-CSF"), or other growth factors.

[0266] Antibodies may also be attached to solid supports, which are particularly useful for immunoassays or purification of the target antigen. Such solid supports include, but are not limited to, glass, cellulose, polyacrylamide, nylon, polystyrene, polyvinyl chloride or polypropylene.

known. See, for example., Arnon et al., "Monoclonal Antibodies For Immunotargeting Of Drugs In Cancer Therapy", in Monoclonal Antibodies And Cancer Therapy, Reisfeld et al. (eds.), pp. 243-56 (Alan R. Liss, Inc. 1985); Hellstrom et al., "Antibodies For Drug Delivery", in Controlled Drug Delivery (2nd Ed.), Robinson et al. (eds.), pp. 623-53 (Marcel Dekker, Inc. 1987); Thorpe, "Antibody Carriers Of Cytotoxic Agents In Cancer Therapy: A Review", in Monoclonal Antibodies '84: Biological And Clinical Applications, Pinchera et al. (eds.), pp. 475-506 (1985); "Analysis, Results, And Future Prospective Of The Therapeutic Use Of Radiolabeled Antibody In Cancer Therapy", in Monoclonal Antibodies For Cancer Detection And Therapy, Baldwin et al. (eds.), pp. 303-16 (Academic Press 1985), and Thorpe et al., "The Preparation And Cytotoxic Properties Of Antibody-Toxin Conjugates", Immunol. Rev. 62:119-58 (1982).

[0268] Alternatively, an antibody can be conjugated to a second antibody to form an antibody heteroconjugate as described by Segal in U.S. Patent No. 4,676,980, which is incorporated herein by reference in its entirety.

[0269] An antibody, with or without a therapeutic moiety conjugated to it, administered alone or in combination with cytotoxic factor(s) and/or cytokine(s) can be used as a therapeutic.

Immunophenotyping

lines and biological samples. Translation products of the genes of the present invention may be useful as cell specific markers, or more specifically as cellular markers that are differentially expressed at various stages of differentiation and/or maturation of particular cell types. Monoclonal antibodies directed against a specific epitope, or combination of epitopes, will allow for the screening of cellular populations expressing the marker. Various techniques can be utilized using monoclonal antibodies to screen for cellular populations expressing the marker(s), and include magnetic separation using antibody-coated magnetic beads, "panning" with antibody attached to a solid matrix (i.e., plate), and flow cytometry (See, e.g., U.S. Patent 5,985,660; and Morrison et al., Cell, 96:737-49 (1999)).

[0271] These techniques allow for the screening of particular populations of cells, such as might be found with hematological malignancies (i.e. minimal residual disease (MRD) in acute leukemic patients) and "non-self" cells in transplantations to prevent Graft-versus-Host Disease (GVHD). Alternatively, these techniques allow for the screening of hematopoietic stem and progenitor cells capable of undergoing proliferation and/or differentiation, as might be found in human umbilical cord blood.

Assays For Antibody Binding

[0272] The antibodies of the invention may be assayed for immunospecific binding by any method known in the art. The immunoassays which can be used include but are not limited to competitive and non-competitive assay systems using techniques such as western blots, radioimmunoassays, ELISA (enzyme linked immunosorbent assay), "sandwich" immunoassays, immunoprecipitation assays, precipitin reactions, gel diffusion

precipitin reactions, immunodiffusion assays, agglutination assays, complement-fixation assays, immunoradiometric assays, fluorescent immunoassays, and protein A immunoassays, to name but a few. Such assays are routine and well known in the art (see, e.g., Ausubel et al, eds, 1994, Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York, which is incorporated by reference herein in its entirety). Exemplary immunoassays are described briefly below (but are not intended by way of limitation).

Immunoprecipitation protocols generally comprise lysing a population of cells [0273] in a lysis buffer such as RIPA buffer (1% NP-40 or Triton X- 100, 1% sodium deoxycholate, 0.1% SDS, 0.15 M NaCl, 0.01 M sodium phosphate at pH 7.2, 1% Trasylol) supplemented with protein phosphatase and/or protease inhibitors (e.g., EDTA, PMSF, aprotinin, sodium vanadate), adding the antibody of interest to the cell lysate, incubating for a period of time (e.g., 1-4 hours) at 4° C, adding protein A and/or protein G sepharose beads to the cell lysate, incubating for about an hour or more at 4° C, washing the beads in lysis buffer and resuspending the beads in SDS/sample buffer. The ability of the antibody of interest to immunoprecipitate a particular antigen can be assessed by, e.g., western blot analysis. One of skill in the art would be knowledgeable as to the parameters that can be modified to increase the binding of the antibody to an antigen and decrease the background (e.g., pre-clearing the cell lysate with sepharose beads). For further discussion regarding immunoprecipitation protocols see, e.g., Ausubel et al., eds., (1994), Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York, section 10.16.1.

[0274] Western blot analysis generally comprises preparing protein samples, electrophoresis of the protein samples in a polyacrylamide gel (e.g., 8%- 20% SDS-PAGE depending on the molecular weight of the antigen), transferring the protein sample from the polyacrylamide gel to a membrane such as nitrocellulose, PVDF or nylon, blocking the membrane in blocking solution (e.g., PBS with 3% BSA or non-fat milk), washing the membrane in washing buffer (e.g., PBS-Tween 20), blocking the membrane with primary antibody (the antibody of interest) diluted in blocking buffer, washing the membrane in washing buffer, blocking the membrane with a secondary antibody (which recognizes the primary antibody, e.g., an anti-human antibody) conjugated to an enzymatic substrate (e.g., horseradish peroxidase or alkaline phosphatase) or radioactive molecule (e.g., 32P or

125I) diluted in blocking buffer, washing the membrane in wash buffer, and detecting the presence of the antigen. One of skill in the art would be knowledgeable as to the parameters that can be modified to increase the signal detected and to reduce the background noise. For further discussion regarding western blot protocols see, e.g., Ausubel et al., eds., (1994), Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York, section 10.8.1.

- ELISAs comprise preparing antigen, coating the well of a 96 well microtiter plate with the antigen, adding the antibody of interest conjugated to a detectable compound such as an enzymatic substrate (e.g., horseradish peroxidase or alkaline phosphatase) to the well and incubating for a period of time, and detecting the presence of the antigen. In ELISAs the antibody of interest does not have to be conjugated to a detectable compound; instead, a second antibody (which recognizes the antibody of interest) conjugated to a detectable compound may be added to the well. Further, instead of coating the well with the antigen, the antibody may be coated to the well. In this case, a second antibody conjugated to a detectable compound may be added following the addition of the antigen of interest to the coated well. One of skill in the art would be knowledgeable as to the parameters that can be modified to increase the signal detected as well as other variations of ELISAs known in the art. For further discussion regarding ELISAs see, e.g., Ausubel et al., eds., (1994), Current Protocols in Molecular Biology, Vol. 1, John Wiley & Sons, Inc., New York, section 11.2.1.
- [0276] The binding affinity of an antibody to an antigen and the off-rate of an antibody-antigen interaction can be determined by competitive binding assays. One example of a competitive binding assay is a radioimmunoassay comprising the incubation of labeled antigen (e.g., 3H or 125I) with the antibody of interest in the presence of increasing amounts of unlabeled antigen, and the detection of the antibody bound to the labeled antigen. The affinity of the antibody of interest for a particular antigen and the binding off-rates can be determined from the data by scatchard plot analysis. Competition with a second antibody can also be determined using radioimmunoassays. In this case, the antigen is incubated with antibody of interest conjugated to a labeled compound (e.g., 3H or 125I) in the presence of increasing amounts of an unlabeled second antibody.
- [0277] Antibodies of the invention may be characterized using immunocytochemisty methods on cells (e.g., mammalian cells, such as CHO cells) transfected with a vector

enabling the expression of an immune/hematopoietic antigen or with vector alone using techniques commonly known in the art. Antibodies that bind immune/hematopoietic antigen transfected cells, but not vector-only transfected cells, are immune/hematopoietic antigen specific.

Therapeutic Uses

[0278] The present invention is further directed to antibody-based therapies which involve administering antibodies of the invention to an animal, preferably a mammal, and most preferably a human, patient for treating one or more of the disclosed diseases. disorders, or conditions. Therapeutic compounds of the invention include, but are not limited to, antibodies of the invention (including fragments, analogs and derivatives thereof as described herein) and nucleic acids encoding antibodies of the invention (including fragments, analogs and derivatives thereof and anti-idiotypic antibodies as described herein). The antibodies of the invention can be used to treat, inhibit or prevent diseases, disorders or conditions associated with aberrant expression and/or activity of a polypeptide of the invention, including, but not limited to, any one or more of the diseases, disorders, or conditions described herein. The treatment and/or prevention of diseases, disorders, or conditions associated with aberrant expression and/or activity of a polypeptide of the invention includes, but is not limited to, alleviating symptoms associated with those diseases, disorders or conditions. Antibodies of the invention may be provided in pharmaceutically acceptable compositions as known in the art or as described herein.

In a specific and preferred embodiment, the present invention is directed to antibody-based therapies which involve administering antibodies of the invention to an animal, preferably a mammal, and most preferably a human, patient for treating one or more of the diseases, disorders, or conditions associated with hematopoiesis or the immune system, including, but not limited to, anemias, autoimmune diseases, immunodeficiencies, allergies, asthma, inflammation and infectious/parasitic diseases. Therapeutic compounds of the invention include, but are not limited to, antibodies of the invention (e.g., antibodies directed to the full length protein expressed on the cell surface of a mammalian cell; antibodies directed to an epitope of an immune/hematopoietic associated polypeptide of the invention (such as, a linear epitope (shown in Table 1A,

column 6) or a conformational epitope), including fragments, analogs and derivatives thereof as described herein) and nucleic acids encoding antibodies of the invention (including fragments, analogs and derivatives thereof and anti-idiotypic antibodies as described herein). The antibodies of the invention can be used to treat, inhibit or prevent diseases, disorders or conditions associated with aberrant expression and/or activity of a polypeptide of the invention, including, but not limited to, any one or more of the diseases, disorders, or conditions associated with hematopoiesis or the immune system described herein. The treatment and/or prevention of diseases, disorders, or conditions of hematopoiesis or the immune system associated with aberrant expression and/or activity of a polypeptide of the invention includes, but is not limited to, alleviating symptoms associated with those diseases, disorders or conditions. Antibodies of the invention may be provided in pharmaceutically acceptable compositions as known in the art or as described herein.

- [0280] A summary of the ways in which the antibodies of the present invention may be used therapeutically includes binding polynucleotides or polypeptides of the present invention locally or systemically in the body or by direct cytotoxicity of the antibody, e.g. as mediated by complement (CDC) or by effector cells (ADCC). Some of these approaches are described in more detail below. Armed with the teachings provided herein, one of ordinary skill in the art will know how to use the antibodies of the present invention for diagnostic, monitoring or therapeutic purposes without undue experimentation.
- [0281] The antibodies of this invention may be advantageously utilized in combination with other monoclonal or chimeric antibodies, or with lymphokines or hematopoietic growth factors (such as, e.g., IL-2, IL-3 and IL-7), for example, which serve to increase the number or activity of effector cells which interact with the antibodies.
- [0282] The antibodies of the invention may be administered alone or in combination with other types of treatments (e.g., radiation therapy, chemotherapy, hormonal therapy, immunotherapy and anti-tumor agents). Generally, administration of products of a species origin or species reactivity (in the case of antibodies) that is the same species as that of the patient is preferred. Thus, in a preferred embodiment, human antibodies, fragments

derivatives, analogs, or nucleic acids, are administered to a human patient for therapy or prophylaxis.

[0283] It is preferred to use high affinity and/or potent *in vivo* inhibiting and/or neutralizing antibodies against polypeptides or polynucleotides of the present invention, fragments or regions thereof, for both immunoassays directed to and therapy of disorders related to polynucleotides or polypeptides, including fragments thereof, of the present invention. Such antibodies, fragments, or regions, will preferably have an affinity for polynucleotides or polypeptides of the invention, including fragments thereof. Preferred binding affinities include those with a dissociation constant or Kd less than 5 X 10⁻² M, 10⁻² M, 5 X 10⁻³ M, 10⁻³ M, 5 X 10⁻⁴ M, 10⁻⁴ M, 5 X 10⁻⁵ M, 10⁻⁵ M, 5 X 10⁻⁶ M, 10⁻⁶ M, 5 X 10⁻⁷ M, 10⁻⁷ M, 5 X 10⁻⁸ M, 10⁻⁸ M, 5 X 10⁻⁹ M, 10⁻⁹ M, 5 X 10⁻¹⁰ M, 10⁻¹⁰ M, 5 X 10⁻¹¹ M, 5 X 10⁻¹² M, 10⁻¹² M, 5 X 10⁻¹³ M, 10⁻¹³ M, 5 X 10⁻¹⁴ M, 10⁻¹⁴ M, 5 X 10⁻¹⁵ M, and 10⁻¹⁵ M.

Gene Therapy

In a specific embodiment, nucleic acids comprising sequences encoding antibodies or functional derivatives thereof, are administered to treat, inhibit or prevent a disease or disorder associated with aberrant expression and/or activity of a polypeptide of the invention, by way of gene therapy. Gene therapy refers to therapy performed by the administration to a subject of an expressed or expressible nucleic acid. In this embodiment of the invention, the nucleic acids produce their encoded protein that mediates a therapeutic effect.

[0285] Any of the methods for gene therapy available in the art can be used according to the present invention. Exemplary methods are described below.

[0286] For general reviews of the methods of gene therapy, see Goldspiel et al., Clinical Pharmacy 12:488-505 (1993); Wu and Wu, Biotherapy 3:87-95 (1991); Tolstoshev, Ann. Rev. Pharmacol. Toxicol. 32:573-596 (1993); Mulligan, Science 260:926-932 (1993); and Morgan and Anderson, Ann. Rev. Biochem. 62:191-217 (1993); May, TIBTECH 11(5):155-215 (1993). Methods commonly known in the art of recombinant DNA technology which can be used are described in Ausubel et al. (eds.), Current Protocols in Molecular Biology, John Wiley & Sons, NY (1993); and Kriegler, Gene Transfer and Expression, A Laboratory Manual, Stockton Press, NY (1990).

In a preferred embodiment, the compound comprises nucleic acid sequences encoding an antibody, said nucleic acid sequences being part of expression vectors that express the antibody or fragments or chimeric proteins or heavy or light chains thereof in a suitable host. In particular, such nucleic acid sequences have promoters operably linked to the antibody coding region, said promoter being inducible or constitutive, and, optionally, tissue-specific. In another particular embodiment, nucleic acid molecules are used in which the antibody coding sequences and any other desired sequences are flanked by regions that promote homologous recombination at a desired site in the genome, thus providing for intrachromosomal expression of the antibody encoding nucleic acids (Koller and Smithies, Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); Zijlstra et al., Nature 342:435-438 (1989). In specific embodiments, the expressed antibody molecule is a single chain antibody; alternatively, the nucleic acid sequences include sequences encoding both the heavy and light chains, or fragments thereof, of the antibody.

[0288] Delivery of the nucleic acids into a patient may be either direct, in which case the patient is directly exposed to the nucleic acid or nucleic acid- carrying vectors, or indirect, in which case, cells are first transformed with the nucleic acids *in vitro*, then transplanted into the patient. These two approaches are known, respectively, as *in vivo* or ex vivo gene therapy.

In a specific embodiment, the nucleic acid sequences are directly administered in vivo, where it is expressed to produce the encoded product. This can be accomplished by any of numerous methods known in the art, e.g., by constructing them as part of an appropriate nucleic acid expression vector and administering it so that they become intracellular, e.g., by infection using defective or attenuated retrovirals or other viral vectors (see U.S. Patent No. 4,980,286), or by direct injection of naked DNA, or by use of microparticle bombardment (e.g., a gene gun; Biolistic, Dupont), or coating with lipids or cell-surface receptors or transfecting agents, encapsulation in liposomes, microparticles, or microcapsules, or by administering them in linkage to a peptide which is known to enter the nucleus, by administering it in linkage to a ligand subject to receptor-mediated endocytosis (see, e.g., Wu and Wu, J. Biol. Chem. 262:4429-4432 (1987)) (which can be used to target cell types specifically expressing the receptors), etc. In another embodiment, nucleic acid-ligand complexes can be formed in which the ligand comprises a fusogenic viral peptide to disrupt endosomes, allowing the nucleic acid to avoid

lysosomal degradation. In yet another embodiment, the nucleic acid can be targeted *in vivo* for cell specific uptake and expression, by targeting a specific receptor (see, e.g., PCT Publications WO 92/06180; WO 92/22635; WO92/20316; WO93/14188, WO 93/20221). Alternatively, the nucleic acid can be introduced intracellularly and incorporated within host cell DNA for expression, by homologous recombination (Koller and Smithies, Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); Zijlstra et al., Nature 342:435-438 (1989)).

In a specific embodiment, viral vectors that contains nucleic acid sequences encoding an antibody of the invention are used. For example, a retroviral vector can be used (see Miller et al., Meth. Enzymol. 217:581-599 (1993)). These retroviral vectors contain the components necessary for the correct packaging of the viral genome and integration into the host cell DNA. The nucleic acid sequences encoding the antibody to be used in gene therapy are cloned into one or more vectors, which facilitates delivery of the gene into a patient. More detail about retroviral vectors can be found in Boesen et al., Biotherapy 6:291-302 (1994), which describes the use of a retroviral vector to deliver the mdr1 gene to hematopoietic stem cells in order to make the stem cells more resistant to chemotherapy. Other references illustrating the use of retroviral vectors in gene therapy are: Clowes et al., J. Clin. Invest. 93:644-651 (1994); Kiem et al., Blood 83:1467-1473 (1994); Salmons and Gunzberg, Human Gene Therapy 4:129-141 (1993); and Grossman and Wilson, Curr. Opin. in Genetics and Devel. 3:110-114 (1993).

Adenoviruses are other viral vectors that can be used in gene therapy. Adenoviruses are especially attractive vehicles for delivering genes to respiratory epithelia. Adenoviruses naturally infect respiratory epithelia where they cause a mild disease. Other targets for adenovirus-based delivery systems are liver, the central nervous system, endothelial cells, and muscle. Adenoviruses have the advantage of being capable of infecting non-dividing cells. Kozarsky and Wilson, Current Opinion in Genetics and Development 3:499-503 (1993) present a review of adenovirus-based gene therapy. Bout et al., Human Gene Therapy 5:3-10 (1994) demonstrated the use of adenovirus vectors to transfer genes to the respiratory epithelia of rhesus monkeys. Other instances of the use of adenoviruses in gene therapy can be found in Rosenfeld et al., Science 252:431-434 (1991); Rosenfeld et al., Cell 68:143- 155 (1992); Mastrangeli et al., J. Clin. Invest.

91:225-234 (1993); PCT Publication WO94/12649; and Wang, et al., Gene Therapy 2:775-783 (1995). In a preferred embodiment, adenovirus vectors are used.

- [0292] Adeno-associated virus (AAV) has also been proposed for use in gene therapy (Walsh et al., Proc. Soc. Exp. Biol. Med. 204:289-300 (1993); U.S. Patent No. 5,436,146).
- [0293] Another approach to gene therapy involves transferring a gene to cells in tissue culture by such methods as electroporation, lipofection, calcium phosphate mediated transfection, or viral infection. Usually, the method of transfer includes the transfer of a selectable marker to the cells. The cells are then placed under selection to isolate those cells that have taken up and are expressing the transferred gene. Those cells are then delivered to a patient.
- In this embodiment, the nucleic acid is introduced into a cell prior to administration *in vivo* of the resulting recombinant cell. Such introduction can be carried out by any method known in the art, including but not limited to transfection, electroporation, microinjection, infection with a viral or bacteriophage vector containing the nucleic acid sequences, cell fusion, chromosome-mediated gene transfer, microcell-mediated gene transfer, spheroplast fusion, etc. Numerous techniques are known in the art for the introduction of foreign genes into cells (see, e.g., Loeffler and Behr, Meth. Enzymol. 217:599-618 (1993); Cohen et al., Meth. Enzymol. 217:618-644 (1993); Cline, Pharmac. Ther. 29:69-92m (1985) and may be used in accordance with the present invention, provided that the necessary developmental and physiological functions of the recipient cells are not disrupted. The technique should provide for the stable transfer of the nucleic acid to the cell, so that the nucleic acid is expressible by the cell and preferably heritable and expressible by its cell progeny.
- [0295] The resulting recombinant cells can be delivered to a patient by various methods known in the art. Recombinant blood cells (e.g., hematopoietic stem or progenitor cells) are preferably administered intravenously. The amount of cells envisioned for use depends on the desired effect, patient state, etc., and can be determined by one skilled in the art.
- [0296] Cells into which a nucleic acid can be introduced for purposes of gene therapy encompass any desired, available cell type, and include but are not limited to epithelial cells, endothelial cells, keratinocytes, fibroblasts, muscle cells, hepatocytes; blood cells

such as T lymphocytes, B lymphocytes, monocytes, macrophages, neutrophils, eosinophils, megakaryocytes, granulocytes; various stem or progenitor cells, in particular hematopoietic stem or progenitor cells, e.g., as obtained from bone marrow, umbilical cord blood, peripheral blood, fetal liver, etc.

[0297] In a preferred embodiment, the cell used for gene therapy is autologous to the patient.

In an embodiment in which recombinant cells are used in gene therapy, nucleic acid sequences encoding an antibody are introduced into the cells such that they are expressible by the cells or their progeny, and the recombinant cells are then administered in vivo for therapeutic effect. In a specific embodiment, stem or progenitor cells are used. Any stem and/or progenitor cells which can be isolated and maintained in vitro can potentially be used in accordance with this embodiment of the present invention (see e.g. PCT Publication WO 94/08598; Stemple and Anderson, Cell 71:973-985 (1992); Rheinwald, Meth. Cell Bio. 21A:229 (1980); and Pittelkow and Scott, Mayo Clinic Proc. 61:771 (1986)).

[0299] In a specific embodiment, the nucleic acid to be introduced for purposes of gene therapy comprises an inducible promoter operably linked to the coding region, such that expression of the nucleic acid is controllable by the presence or absence of an appropriate inducer of transcription.

Demonstration of Therapeutic or Prophylactic Activity

tested *in vitro*, and then *in vivo* for the desired therapeutic or prophylactic activity, prior to use in humans. For example, *in vitro* assays to demonstrate the therapeutic or prophylactic utility of a compound or pharmaceutical composition include, the effect of a compound on a cell line or a patient tissue sample. The effect of the compound or composition on the cell line and/or tissue sample can be determined utilizing techniques known to those of skill in the art including, but not limited to, rosette formation assays and cell lysis assays. In accordance with the invention, *in vitro* assays which can be used to determine whether administration of a specific compound is indicated, include *in vitro* cell culture assays in which a patient tissue sample is grown in culture, and exposed to or

otherwise administered a compound, and the effect of such compound upon the tissue sample is observed.

Therapeutic/Prophylactic Administration and Composition

[0301] The invention provides methods of treatment, inhibition and prophylaxis by administration to a subject of an effective amount of a compound or pharmaceutical composition of the invention, preferably a polypeptide or antibody of the invention. In a preferred embodiment, the compound is substantially purified (e.g., substantially free from substances that limit its effect or produce undesired side-effects). The subject is preferably an animal, including but not limited to animals such as cows, pigs, horses, chickens, cats, dogs, etc., and is preferably a mammal, and most preferably human.

[0302] Formulations and methods of administration that can be employed when the compound comprises a nucleic acid or an immunoglobulin are described above; additional appropriate formulations and routes of administration can be selected from among those described herein below.

[0303] Various delivery systems are known and can be used to administer a compound of the invention, e.g., encapsulation in liposomes, microparticles, microcapsules, recombinant cells capable of expressing the compound, receptor-mediated endocytosis (see, e.g., Wu and Wu, J. Biol. Chem. 262:4429-4432 (1987)), construction of a nucleic acid as part of a retroviral or other vector, etc. Methods of introduction include but are not limited to intradermal, intramuscular, intraperitoneal, intravenous, subcutaneous, intranasal, epidural, and oral routes. The compounds or compositions may be administered by any convenient route, for example by infusion or bolus injection, by absorption through epithelial or mucocutaneous linings (e.g., oral mucosa, rectal and intestinal mucosa, etc.) and may be administered together with other biologically active agents. Administration can be systemic or local. In addition, it may be desirable to introduce the pharmaceutical compounds or compositions of the invention into the central nervous system by any suitable route, including intraventricular and intrathecal injection; intraventricular injection may be facilitated by an intraventricular catheter, for example, attached to a reservoir, such as an Ommaya reservoir. Pulmonary administration can also be employed, e.g., by use of an inhaler or nebulizer, and formulation with an aerosolizing agent.

[0304] In a specific embodiment, it may be desirable to administer the pharmaceutical compounds or compositions of the invention locally to the area in need of treatment; this may be achieved by, for example, and not by way of limitation, local infusion during surgery, topical application, e.g., in conjunction with a wound dressing after surgery, by injection, by means of a catheter, by means of a suppository, or by means of an implant, said implant being of a porous, non-porous, or gelatinous material, including membranes, such as sialastic membranes, or fibers. Preferably, when administering a protein, including an antibody, of the invention, care must be taken to use materials to which the protein does not absorb.

- [0305] In another embodiment, the compound or composition can be delivered in a vesicle, in particular a liposome (see Langer, Science 249:1527-1533 (1990); Treat et al., in Liposomes in the Therapy of Infectious Disease and Cancer, Lopez-Berestein and Fidler (eds.), Liss, New York, pp. 353- 365 (1989); Lopez-Berestein, ibid., pp. 317-327; see generally ibid.)
- In yet another embodiment, the compound or composition can be delivered in a controlled release system. In one embodiment, a pump may be used (see Langer, supra; Sefton, CRC Crit. Ref. Biomed. Eng. 14:201 (1987); Buchwald et al., Surgery 88:507 (1980); Saudek et al., N. Engl. J. Med. 321:574 (1989)). In another embodiment, polymeric materials can be used (see Medical Applications of Controlled Release, Langer and Wise (eds.), CRC Pres., Boca Raton, Florida (1974); Controlled Drug Bioavailability, Drug Product Design and Performance, Smolen and Ball (eds.), Wiley, New York (1984); Ranger and Peppas, J., Macromol. Sci. Rev. Macromol. Chem. 23:61 (1983); see also Levy et al., Science 228:190 (1985); During et al., Ann. Neurol. 25:351 (1989); Howard et al., J.Neurosurg. 71:105 (1989)). In yet another embodiment, a controlled release system can be placed in proximity of the therapeutic target, e.g., the brain, thus requiring only a fraction of the systemic dose (see, e.g., Goodson, in Medical Applications of Controlled Release, supra, vol. 2, pp. 115-138 (1984)).
- [0307] Other controlled release systems are discussed in the review by Langer (Science 249:1527-1533 (1990)).
- [0308] In a specific embodiment where the compound of the invention is a nucleic acid encoding a protein, the nucleic acid can be administered *in vivo* to promote expression of its encoded protein, by constructing it as part of an appropriate nucleic acid

expression vector and administering it so that it becomes intracellular, e.g., by use of a retroviral vector (see U.S. Patent No. 4,980,286), or by direct injection, or by use of microparticle bombardment (e.g., a gene gun; Biolistic, Dupont), or coating with lipids or cell-surface receptors or transfecting agents, or by administering it in linkage to a homeobox-like peptide which is known to enter the nucleus (see e.g., Joliot et al., Proc. Natl. Acad. Sci. USA 88:1864-1868 (1991)), etc. Alternatively, a nucleic acid can be introduced intracellularly and incorporated within host cell DNA for expression, by homologous recombination.

[0309] The present invention also provides pharmaceutical compositions. Such compositions comprise a therapeutically effective amount of a compound, and a pharmaceutically acceptable carrier. In a specific embodiment, the term "pharmaceutically acceptable" means approved by a regulatory agency of the Federal or a state government or listed in the U.S. Pharmacopeia or other generally recognized pharmacopeia for use in animals, and more particularly in humans. The term "carrier" refers to a diluent, adjuvant, excipient, or vehicle with which the therapeutic is administered. Such pharmaceutical carriers can be sterile liquids, such as water and oils, including those of petroleum, animal, vegetable or synthetic origin, such as peanut oil, soybean oil, mineral oil, sesame oil and the like. Water is a preferred carrier when the pharmaceutical composition is administered intravenously. Saline solutions and aqueous dextrose and glycerol solutions can also be employed as liquid carriers, particularly for injectable solutions. Suitable pharmaceutical excipients include starch, glucose, lactose, sucrose, gelatin, malt, rice, flour, chalk, silica gel, sodium stearate, glycerol monostearate, talc, sodium chloride, dried skim milk, glycerol, propylene, glycol, water, ethanol and the The composition, if desired, can also contain minor amounts of wetting or emulsifying agents, or pH buffering agents. These compositions can take the form of solutions, suspensions, emulsion, tablets, pills, capsules, powders, sustained-release formulations and the like. The composition can be formulated as a suppository, with traditional binders and carriers such as triglycerides. Oral formulation can include standard carriers such as pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharine, cellulose, magnesium carbonate, etc. Examples of suitable pharmaceutical carriers are described in "Remington's Pharmaceutical Sciences" by E.W. Martin. Such compositions will contain a therapeutically effective amount of the

compound, preferably in purified form, together with a suitable amount of carrier so as to provide the form for proper administration to the patient. The formulation should suit the mode of administration.

- In a preferred embodiment, the composition is formulated in accordance with routine procedures as a pharmaceutical composition adapted for intravenous administration to human beings. Typically, compositions for intravenous administration are solutions in sterile isotonic aqueous buffer. Where necessary, the composition may also include a solubilizing agent and a local anesthetic such as lignocaine to ease pain at the site of the injection. Generally, the ingredients are supplied either separately or mixed together in unit dosage form, for example, as a dry lyophilized powder or water free concentrate in a hermetically sealed container such as an ampoule or sachette indicating the quantity of active agent. Where the composition is to be administered by infusion, it can be dispensed with an infusion bottle containing sterile pharmaceutical grade water or saline. Where the composition is administered by injection, an ampoule of sterile water for injection or saline can be provided so that the ingredients may be mixed prior to administration.
- [0311] The compounds of the invention can be formulated as neutral or salt forms. Pharmaceutically acceptable salts include those formed with anions such as those derived from hydrochloric, phosphoric, acetic, oxalic, tartaric acids, etc., and those formed with cations such as those derived from sodium, potassium, ammonium, calcium, ferric hydroxides, isopropylamine, triethylamine, 2-ethylamino ethanol, histidine, procaine, etc.
- [0312] The amount of the compound of the invention which will be effective in the treatment, inhibition and prevention of a disease or disorder associated with aberrant expression and/or activity of a polypeptide of the invention can be determined by standard clinical techniques. In addition, in vitro assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also depend on the route of administration, and the seriousness of the disease or disorder, and should be decided according to the judgment of the practitioner and each patient's circumstances. Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems.
- [0313] For antibodies, the dosage administered to a patient is typically 0.1 mg/kg to 100 mg/kg of the patient's body weight. Preferably, the dosage administered to a patient

is between 0.1 mg/kg and 20 mg/kg of the patient's body weight, more preferably 1 mg/kg to 10 mg/kg of the patient's body weight. Generally, human antibodies have a longer half-life within the human body than antibodies from other species due to the immune response to the foreign polypeptides. Thus, lower dosages of human antibodies and less frequent administration is often possible. Further, the dosage and frequency of administration of antibodies of the invention may be reduced by enhancing uptake and tissue penetration (e.g., into the brain) of the antibodies by modifications such as, for example, lipidation.

[0314] The invention also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the pharmaceutical compositions of the invention. Optionally associated with such container(s) can be a notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of pharmaceuticals or biological products, which notice reflects approval by the agency of manufacture, use or sale for human administration.

Diagnosis and Imaging

[0315] Labeled antibodies, and derivatives and analogs thereof, which specifically bind to a polypeptide of interest can be used for diagnostic purposes to detect, diagnose, or monitor diseases, disorders, and/or conditions associated with the aberrant expression and/or activity of a polypeptide of the invention. The invention provides for the detection of aberrant expression of a polypeptide of interest, comprising (a) assaying the expression of the polypeptide of interest in cells or body fluid of an individual using one or more antibodies specific to the polypeptide interest and (b) comparing the level of gene expression with a standard gene expression level, whereby an increase or decrease in the assayed polypeptide gene expression level compared to the standard expression level is indicative of aberrant expression.

[0316] The invention provides a diagnostic assay for diagnosing an immune/hematopoietic-related disorder, comprising (a) assaying the expression of the polypeptide of interest in cells or body fluid of an individual using one or more antibodies specific to the polypeptide interest and (b) comparing the level of gene expression with a standard gene expression level, whereby an increase or decrease in the assayed polypeptide gene expression level compared to the standard expression level is indicative

of a particular disorder. With respect to cancer, the presence of a relatively high amount of transcript in biopsied tissue from an individual may indicate a predisposition for the development of the disease, or may provide a means for detecting the disease prior to the appearance of actual clinical symptoms. A more definitive diagnosis of this type may allow health professionals to employ preventative measures or aggressive treatment earlier thereby preventing the development or further progression of the cancer.

[0317] Antibodies of the invention can be used to assay protein levels in a biological sample using classical immunohistological methods known to those of skill in the art (e.g., see Jalkanen et al., J. Cell. Biol. 101:976-985 (1985); Jalkanen et al., J. Cell. Biol. 105:3087-3096 (1987)). Other antibody-based methods useful for detecting protein gene expression include immunoassays, such as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). Suitable antibody assay labels are known in the art and include enzyme labels, such as, glucose oxidase; radioisotopes, such as iodine (125I, 121I), carbon (14C), sulfur (35S), tritium (3H), indium (112In), and technetium (99Tc); luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

One facet of the invention is the detection and diagnosis of a disease or [0318] disorder associated with aberrant expression of a polypeptide of interest in an animal, preferably a mammal and most preferably a human. A preferred embodiment of the invention is the detection and diagnosis of a disease or disorder of hematopoiesis or the immune system associated with aberrant expression of an immune/hematopoietic antigen in an animal, preferably a mammal and most preferably a human. In one embodiment, diagnosis comprises: a) administering (for example, parenterally, subcutaneously, or intraperitoneally) to a subject an effective amount of a labeled molecule which specifically binds to the polypeptide of interest; b) waiting for a time interval following the administering for permitting the labeled molecule to preferentially concentrate at sites in the subject where the polypeptide is expressed (and for unbound labeled molecule to be cleared to background level); c) determining background level; and d) detecting the labeled molecule in the subject, such that detection of labeled molecule above the background level indicates that the subject has a particular disease or disorder associated with aberrant expression of the polypeptide of interest. Background level can be

determined by various methods including, comparing the amount of labeled molecule detected to a standard value previously determined for a particular system.

- [0319] It will be understood in the art that the size of the subject and the imaging system used will determine the quantity of imaging moiety needed to produce diagnostic images. In the case of a radioisotope moiety, for a human subject, the quantity of radioactivity injected will normally range from about 5 to 20 millicuries of 99mTc. The labeled antibody or antibody fragment will then preferentially accumulate at the location of cells which contain the specific protein. *In vivo* tumor imaging is described in S.W. Burchiel et al., "Immunopharmacokinetics of Radiolabeled Antibodies and Their Fragments." (Chapter 13 in Tumor Imaging: The Radiochemical Detection of Cancer, S.W. Burchiel and B. A. Rhodes, eds., Masson Publishing Inc. (1982)).
- [0320] Depending on several variables, including the type of label used and the mode of administration, the time interval following the administration for permitting the labeled molecule to preferentially concentrate at sites in the subject and for unbound labeled molecule to be cleared to background level is 6 to 48 hours or 6 to 24 hours or 6 to 12 hours. In another embodiment the time interval following administration is 5 to 20 days or 5 to 10 days.
- [0321] In an embodiment, monitoring of the disease or disorder is carried out by repeating the method for diagnosing the disease or disorder, for example, one month after initial diagnosis, six months after initial diagnosis, one year after initial diagnosis, etc.
- [0322] Presence of the labeled molecule can be detected in the patient using methods known in the art for *in vivo* scanning. These methods depend upon the type of label used. Skilled artisans will be able to determine the appropriate method for detecting a particular label. Methods and devices that may be used in the diagnostic methods of the invention include, but are not limited to, computed tomography (CT), whole body scan such as position emission tomography (PET), magnetic resonance imaging (MRI), and sonography.
- [0323] In a specific embodiment, the molecule is labeled with a radioisotope and is detected in the patient using a radiation responsive surgical instrument (Thurston et al., U.S. Patent No. 5,441,050). In another embodiment, the molecule is labeled with a fluorescent compound and is detected in the patient using a fluorescence responsive scanning instrument. In another embodiment, the molecule is labeled with a positron

emitting metal and is detected in the patent using positron emission-tomography. In yet another embodiment, the molecule is labeled with a paramagnetic label and is detected in a patient using magnetic resonance imaging (MRI).

Kits

In one embodiment, a kit comprises an antibody of the invention, preferably a purified antibody, in one or more containers. In a specific embodiment, the kits of the present invention contain a substantially isolated polypeptide comprising an epitope which is specifically immunoreactive with an antibody included in the kit. Preferably, the kits of the present invention further comprise a control antibody which does not react with the polypeptide of interest. In another specific embodiment, the kits of the present invention contain a means for detecting the binding of an antibody to a polypeptide of interest (e.g., the antibody may be conjugated to a detectable substrate such as a fluorescent compound, an enzymatic substrate, a radioactive compound or a luminescent compound, or a second antibody which recognizes the first antibody may be conjugated to a detectable substrate).

[0325] In another specific embodiment of the present invention, the kit is a diagnostic kit for use in screening serum containing antibodies specific against proliferative and/or cancerous polynucleotides and polypeptides. Such a kit may include a control antibody that does not react with the polypeptide of interest. Such a kit may include a substantially isolated polypeptide antigen comprising an epitope which is specifically immunoreactive with at least one anti-polypeptide antigen antibody. Further, such a kit includes means for detecting the binding of said antibody to the antigen (e.g., the antibody may be conjugated to a fluorescent compound such as fluorescein or rhodamine which can be detected by flow cytometry). In specific embodiments, the kit may include a recombinantly produced or chemically synthesized polypeptide antigen. The polypeptide antigen of the kit may also be attached to a solid support.

[0326] In a more specific embodiment the detecting means of the above-described kit includes a solid support to which said polypeptide antigen is attached. Such a kit may also include a non-attached reporter-labeled anti-human antibody. In this embodiment, binding of the antibody to the polypeptide antigen can be detected by binding of the said reporter-labeled antibody.

In an additional embodiment, the invention includes a diagnostic kit for use in screening serum containing antigens of the polypeptide of the invention. The diagnostic kit includes a substantially isolated antibody specifically immunoreactive with polypeptide or polynucleotide antigens, and means for detecting the binding of the polynucleotide or polypeptide antigen to the antibody. In one embodiment, the antibody is attached to a solid support. In a specific embodiment, the antibody may be a monoclonal antibody. The detecting means of the kit may include a second, labeled monoclonal antibody. Alternatively, or in addition, the detecting means may include a labeled, competing antigen.

- In one diagnostic configuration, test serum is reacted with a solid phase reagent having a surface-bound antigen obtained by the methods of the present invention. After binding with specific antigen antibody to the reagent and removing unbound serum components by washing, the reagent is reacted with reporter-labeled anti-human antibody to bind reporter to the reagent in proportion to the amount of bound anti-antigen antibody on the solid support. The reagent is again washed to remove unbound labeled antibody, and the amount of reporter associated with the reagent is determined. Typically, the reporter is an enzyme which is detected by incubating the solid phase in the presence of a suitable fluorometric, luminescent or colorimetric substrate (Sigma, St. Louis, MO).
- [0329] The solid surface reagent in the above assay is prepared by known techniques for attaching protein material to solid support material, such as polymeric beads, dip sticks, 96-well plate or filter material. These attachment methods generally include non-specific adsorption of the protein to the support or covalent attachment of the protein, typically through a free amine group, to a chemically reactive group on the solid support, such as an activated carboxyl, hydroxyl, or aldehyde group. Alternatively, streptavidin coated plates can be used in conjunction with biotinylated antigen(s).
- [0330] Thus, the invention provides an assay system or kit for carrying out this diagnostic method. The kit generally includes a support with surface-bound recombinant antigens, and a reporter-labeled anti-human antibody for detecting surface-bound anti-antigen antibody.

Uses of the Polynucleotides

[0331] Each of the polynucleotides identified herein can be used in numerous ways as reagents. The following description should be considered exemplary and utilizes known techniques.

- [0332] The polynucleotides of the present invention are useful for chromosome identification. There exists an ongoing need to identify new chromosome markers, since few chromosome marking reagents, based on actual sequence data (repeat polymorphisms), are presently available. Each sequence is specifically targeted to and can hybridize with a particular location on an individual human chromosome, thus each polynucleotide of the present invention can routinely be used as a chromosome marker using techniques known in the art. Table 1A, column 8 provides the chromosome location of some of the polynucleotides of the invention.
- [0333] Briefly, sequences can be mapped to chromosomes by preparing PCR primers (preferably at least 15 bp (e.g., 15-25 bp) from the sequences shown in SEQ ID NO:X. Primers can optionally be selected using computer analysis so that primers do not span more than one predicted exon in the genomic DNA. These primers are then used for PCR screening of somatic cell hybrids containing individual human chromosomes. Only those hybrids containing the human gene corresponding to SEQ ID NO:X will yield an amplified fragment.
- [0334] Similarly, somatic hybrids provide a rapid method of PCR mapping the polynucleotides to particular chromosomes. Three or more clones can be assigned per day using a single thermal cycler. Moreover, sublocalization of the polynucleotides can be achieved with panels of specific chromosome fragments. Other gene mapping strategies that can be used include in situ hybridization, prescreening with labeled flow-sorted chromosomes, preselection by hybridization to construct chromosome specific-cDNA libraries, and computer mapping techniques (See, e.g., Shuler, Trends Biotechnol 16:456-459 (1998) which is hereby incorporated by reference in its entirety).
- [0335] Precise chromosomal location of the polynucleotides can also be achieved using fluorescence in situ hybridization (FISH) of a metaphase chromosomal spread. This technique uses polynucleotides as short as 500 or 600 bases; however, polynucleotides 2,000-4,000 bp are preferred. For a review of this technique, see Verma et al., "Human Chromosomes: a Manual of Basic Techniques," Pergamon Press, New York (1988).

[0336] For chromosome mapping, the polynucleotides can be used individually (to mark a single chromosome or a single site on that chromosome) or in panels (for marking multiple sites and/or multiple chromosomes).

- [0337] Thus, the present invention also provides a method for chromosomal localization which involves (a) preparing PCR primers from the polynucleotide sequences in Table 1A and/or Table 2 and SEQ ID NO:X and (b) screening somatic cell hybrids containing individual chromosomes.
- [0338] The polynucleotides of the present invention would likewise be useful for radiation hybrid mapping, HAPPY mapping, and long range restriction mapping. For a review of these techniques and others known in the art, see, e.g. Dear, "Genome Mapping: A Practical Approach," IRL Press at Oxford University Press, London (1997); Aydin, J. Mol. Med. 77:691-694 (1999); Hacia et al., Mol. Psychiatry 3:483-492 (1998); Herrick et al., Chromosome Res. 7:409-423 (1999); Hamilton et al., Methods Cell Biol. 62:265-280 (2000); and/or Ott, J. Hered. 90:68-70 (1999), each of which is hereby incorporated by reference in its entirety.
- [0339] Once a polynucleotide has been mapped to a precise chromosomal location, the physical position of the polynucleotide can be used in linkage analysis. Linkage analysis establishes coinheritance between a chromosomal location and presentation of a particular disease. (Disease mapping data are found, for example, in V. McKusick, Mendelian Inheritance in Man (available on line through Johns Hopkins University Welch Medical Library).) Column 9 of Table 1A provides an OMIM reference identification number of diseases associated with the cytologic band disclosed in column 8 of Table 1A, as determined using techniques described herein and by reference to Table 5. Assuming 1 megabase mapping resolution and one gene per 20 kb, a cDNA precisely localized to a chromosomal region associated with the disease could be one of 50-500 potential causative genes.
- [0340] Thus, once coinheritance is established, differences in a polynucleotide of the invention and the corresponding gene between affected and unaffected individuals can be examined. First, visible structural alterations in the chromosomes, such as deletions or translocations, are examined in chromosome spreads or by PCR. If no structural alterations exist, the presence of point mutations are ascertained. Mutations observed in some or all affected individuals, but not in normal individuals, indicates that the mutations

may cause the disease. However, complete sequencing of the polypeptide and the corresponding gene from several normal individuals is required to distinguish the mutation from a polymorphism. If a new polymorphism is identified, this polymorphic polypeptide can be used for further linkage analysis.

- [0341] Furthermore, increased or decreased expression of the gene in affected individuals as compared to unaffected individuals can be assessed using the polynucleotides of the invention. Any of these alterations (altered expression, chromosomal rearrangement, or mutation) can be used as a diagnostic or prognostic marker. Diagnostic and prognostic methods, kits and reagents encompassed by the present invention are briefly described below and more thoroughly elsewhere herein (see e.g., the sections labeled "Antibodies", "Diagnostic Assays", and "Methods for Detecting Immune/Hematopoietic-Related Disease, Including Cancer").
- [0342] Thus, the invention also provides a diagnostic method useful during diagnosis of a disorder, involving measuring the expression level of polynucleotides of the present invention in cells or body fluid from an individual and comparing the measured gene expression level with a standard level of polynucleotide expression level, whereby an increase or decrease in the gene expression level compared to the standard is indicative of a disorder. Additional non-limiting examples of diagnostic methods encompassed by the present invention are more thoroughly described elsewhere herein (see, e.g., Example 12).
- [0343] In still another embodiment, the invention includes a kit for analyzing samples for the presence of proliferative and/or cancerous polynucleotides derived from a test subject, as further described herein. In a general embodiment, the kit includes at least one polynucleotide probe containing a nucleotide sequence that will specifically hybridize with a polynucleotide of the invention and a suitable container. In a specific embodiment, the kit includes two polynucleotide probes defining an internal region of the polynucleotide of the invention, where each probe has one strand containing a 31'mer-end internal to the region. In a further embodiment, the probes may be useful as primers for polymerase chain reaction amplification.
- [0344] Where a diagnosis of a related disorder, including, for example, diagnosis of a tumor, has already been made according to conventional methods, the present invention is useful as a prognostic indicator, whereby patients exhibiting enhanced or depressed

polynucleotide of the invention expression will experience a worse clinical outcome relative to patients expressing the gene at a level nearer the standard level.

[0345] By "measuring the expression level of polynucleotides of the invention" is intended qualitatively or quantitatively measuring or estimating the level of the polypeptide of the invention or the level of the mRNA encoding the polypeptide of the invention in a first biological sample either directly (e.g., by determining or estimating absolute protein level or mRNA level) or relatively (e.g., by comparing to the polypeptide level or mRNA level in a second biological sample). Preferably, the polypeptide level or mRNA level in the first biological sample is measured or estimated and compared to a standard polypeptide level or mRNA level, the standard being taken from a second biological sample obtained from an individual not having the related disorder or being determined by averaging levels from a population of individuals not having a related disorder. As will be appreciated in the art, once a standard polypeptide level or mRNA level is known, it can be used repeatedly as a standard for comparison.

[0346] By "biological sample" is intended any biological sample obtained from an individual, body fluid, cell line, tissue culture, or other source which contains polypeptide of the present invention or the corresponding mRNA. As indicated, biological samples include body fluids (such as semen, lymph, vaginal pool, sera, plasma, urine, synovial fluid and spinal fluid) which contain the polypeptide of the present invention, and tissue sources found to express the polypeptide of the present invention. Methods for obtaining tissue biopsies and body fluids from mammals are well known in the art. Where the biological sample is to include mRNA, a tissue biopsy is the preferred source.

[0347] The method(s) provided above may preferably be applied in a diagnostic method and/or kits in which polynucleotides and/or polypeptides of the invention are attached to a solid support. In one exemplary method, the support may be a "gene chip" or a "biological chip" as described in U.S. Patents 5,837,832, 5,874,219, and 5,856,174. Further, such a gene chip with polynucleotides of the invention attached may be used to identify polymorphisms between the isolated polynucleotide sequences of the invention, with polynucleotides isolated from a test subject. The knowledge of such polymorphisms (i.e., their location, as well as, their existence) would be beneficial in identifying disease loci for many disorders, such as for example, in neural disorders, immune system disorders, muscular disorders, reproductive disorders, gastrointestinal disorders,

pulmonary disorders, digestive disorders, cardiovascular disorders, renal disorders, proliferative disorders, and/or cancerous diseases and conditions. Such a method is described in U.S. Patents 5,858,659 and 5,856,104. The U.S. Patents referenced *supra* are hereby incorporated by reference in their entirety herein.

[0348] The present invention encompasses polynucleotides of the present invention that are chemically synthesized, or reproduced as peptide nucleic acids (PNA), or according to other methods known in the art. The use of PNAs would serve as the preferred form if the polynucleotides of the invention are incorporated onto a solid support, or gene chip. For the purposes of the present invention, a peptide nucleic acid (PNA) is a polyamide type of DNA analog and the monomeric units for adenine, guanine, thymine and cytosine are available commercially (Perceptive Biosystems). Certain components of DNA, such as phosphorus, phosphorus oxides, or deoxyribose derivatives, are not present in PNAs. As disclosed by Nielsen et al., Science 254:1497 (1991); and Egholm et al., Nature 365:666 (1993), PNAs bind specifically and tightly to complementary DNA strands and are not degraded by nucleases. In fact, PNA binds more strongly to DNA than DNA itself does. This is probably because there is no electrostatic repulsion between the two strands, and also the polyamide backbone is more flexible. Because of this, PNA/DNA duplexes bind under a wider range of stringency conditions than DNA/DNA duplexes, making it easier to perform multiplex hybridization. Smaller probes can be used than with DNA due to the strong binding. In addition, it is more likely that single base mismatches can be determined with PNA/DNA hybridization because a single mismatch in a PNA/DNA 15-mer lowers the melting point (T.sub.m) by 8°-20° C, vs. 4°-16° C for the DNA/DNA 15-mer duplex. Also, the absence of charge groups in PNA means that hybridization can be done at low ionic strengths and reduce possible interference by salt during the analysis.

[0349] The compounds of the present invention have uses which include, but are not limited to, detecting cancer in mammals. In particular the invention is useful during diagnosis of pathological cell proliferative neoplasias which include, but are not limited to: acute myelogenous leukemias including acute monocytic leukemia, acute myeloblastic leukemia, acute promyelocytic leukemia, acute myelomonocytic leukemia, acute erythroleukemia, acute megakaryocytic leukemia, and acute undifferentiated leukemia, etc.; and chronic myelogenous leukemias including chronic myelomonocytic leukemia,

chronic granulocytic leukemia, etc. Preferred mammals include monkeys, apes, cats, dogs, cows, pigs, horses, rabbits and humans. Particularly preferred are humans.

The compounds of the present invention have preferred uses which include, but are not limited to, detecting cancer of hematopoietic cells in mammals. In particular, the invention is useful during diagnosis of pathological cell proliferative neoplasias which include, but are not limited to: leukemia, acute myelogenous leukemia, chronic myelogeneous leukemia, Hodgkin's disease, non-Hodgkin's lymphoma, acute lymphocytic anemia (ALL) Chronic lymphocyte leukemia, plasmacytomas, multiple myeloma, and Burkitt's lymphoma. Preferred mammals include monkeys, apes, cats, dogs, cows, pigs, horses, rabbits and humans. Particularly preferred are humans.

[0351] Pathological cell proliferative disorders are often associated with inappropriate activation of proto-oncogenes. (Gelmann, E. P. et al., "The Etiology of Acute Leukemia: Molecular Genetics and Viral Oncology," in Neoplastic Diseases of the Blood, Vol 1., Wiernik, P. H. et al. eds., 161-182 (1985)). Neoplasias are now believed to result from the qualitative alteration of a normal cellular gene product, or from the quantitative modification of gene expression by insertion into the chromosome of a viral sequence, by chromosomal translocation of a gene to a more actively transcribed region, or by some other mechanism. (Gelmann et al., *supra*) It is likely that mutated or altered expression of specific genes is involved in the pathogenesis of some leukemias, among other tissues and cell types. (Gelmann et al., *supra*) Indeed, the human counterparts of the oncogenes involved in some animal neoplasias have been amplified or translocated in some cases of human leukemia and carcinoma. (Gelmann et al., *supra*)

[0352] For example, c-myc expression is highly amplified in the non-lymphocytic leukemia cell line HL-60. When HL-60 cells are chemically induced to stop proliferation, the level of c-myc is found to be downregulated. (International Publication Number WO 91/15580). However, it has been shown that exposure of HL-60 cells to a DNA construct that is complementary to the 5' end of c-myc or c-myb blocks translation of the corresponding mRNAs which downregulates expression of the c-myc or c-myb proteins and causes arrest of cell proliferation and differentiation of the treated cells. (International Publication Number WO 91/15580; Wickstrom et al., Proc. Natl. Acad. Sci. 85:1028 (1988); Anfossi et al., Proc. Natl. Acad. Sci. 86:3379 (1989)). However, the skilled artisan would appreciate the present invention's usefulness is not be limited to treatment,

prevention, diagnosis and/or prognosis, of proliferative disorders of cells and tissues of hematopoietic origin, in light of the numerous cells and cell types of varying origins which are known to exhibit proliferative phenotypes. In preferred embodiments, the compounds and/or methods of the invention are used to treat, prevent, diagnose, and/or prognose, proliferative disorders of hematopoietic cells and tissues as well as cells and tissues of the immune system.

[0353] In addition to the foregoing, a polynucleotide of the present invention can be used to control gene expression through triple helix formation or through antisense DNA or RNA. Antisense techniques are discussed, for example, in Okano, J. Neurochem. 56: 560 (1991); "Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988). Triple helix formation is discussed in, for instance Lee et al., Nucleic Acids Research 6: 3073 (1979); Cooney et al., Science 241: 456 (1988); and Dervan et al., Science 251: 1360 (1991). Both methods rely on binding of the polynucleotide to a complementary DNA or RNA. For these techniques, preferred polynucleotides are usually oligonucleotides 20 to 40 bases in length and complementary to either the region of the gene involved in transcription (triple helix - see Lee et al., Nucl. Acids Res. 6:3073 (1979); Cooney et al., Science 241:456 (1988); and Dervan et al., Science 251:1360 (1991)) or to the mRNA itself (antisense - Okano, J. Neurochem. 56:560 (1991); Oligodeoxy-nucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988).) Triple helix formation optimally results in a shut-off of RNA transcription from DNA, while antisense RNA hybridization blocks translation of an mRNA molecule into polypeptide. The oligonucleotide described above can also be delivered to cells such that the antisense RNA or DNA may be expressed in vivo to inhibit production of polypeptide of the present invention antigens. Both techniques are effective in model systems, and the information disclosed herein can be used to design antisense or triple helix polynucleotides in an effort to treat disease, and in particular, for the treatment of proliferative diseases and/or conditions. Non-limiting antisense and triple helix methods encompassed by the present invention are more thoroughly described elsewhere herein (see, e.g., the section labeled "Antisense and Ribozyme (Antagonists)").

[0354] Polynucleotides of the present invention are also useful in gene therapy. One goal of gene therapy is to insert a normal gene into an organism having a defective gene, in an effort to correct the genetic defect. The polynucleotides disclosed in the present

invention offer a means of targeting such genetic defects in a highly accurate manner. Another goal is to insert a new gene that was not present in the host genome, thereby producing a new trait in the host cell. Additional non-limiting examples of gene therapy methods encompassed by the present invention are more thoroughly described elsewhere herein (see, e.g., the sections labeled "Gene Therapy Methods" and Examples 16, 17 and 18).

[0355] The polynucleotides are also useful for identifying individuals from minute biological samples. The United States military, for example, is considering the use of restriction fragment length polymorphism (RFLP) for identification of its personnel. In this technique, an individual's genomic DNA is digested with one or more restriction enzymes, and probed on a Southern blot to yield unique bands for identifying personnel. This method does not suffer from the current limitations of "Dog Tags" which can be lost, switched, or stolen, making positive identification difficult. The polynucleotides of the present invention can be used as additional DNA markers for RFLP.

[0356] The polynucleotides of the present invention can also be used as an alternative to RFLP, by determining the actual base-by-base DNA sequence of selected portions of an individual's genome. These sequences can be used to prepare PCR primers for amplifying and isolating such selected DNA, which can then be sequenced. Using this technique, individuals can be identified because each individual will have a unique set of DNA sequences. Once an unique ID database is established for an individual, positive identification of that individual, living or dead, can be made from extremely small tissue samples.

[0357] Forensic biology also benefits from using DNA-based identification techniques as disclosed herein. DNA sequences taken from very small biological samples such as tissues, e.g., hair or skin, or body fluids, e.g., blood, saliva, semen, synovial fluid, amniotic fluid, breast milk, lymph, pulmonary sputum or surfactant, urine, fecal matter, etc., can be amplified using PCR. In one prior art technique, gene sequences amplified from polymorphic loci, such as DQa class II HLA gene, are used in forensic biology to identify individuals. (Erlich, H., PCR Technology, Freeman and Co. (1992).) Once these specific polymorphic loci are amplified, they are digested with one or more restriction enzymes, yielding an identifying set of bands on a Southern blot probed with DNA

corresponding to the DQa class II HLA gene. Similarly, polynucleotides of the present invention can be used as polymorphic markers for forensic purposes.

[0358] There is also a need for reagents capable of identifying the source of a particular tissue. Such need arises, for example, in forensics when presented with tissue of unknown origin. Appropriate reagents can comprise, for example, DNA probes or primers prepared from the sequences of the present invention, specific to tissues, including but not limited to, those sequences referred to in Table 1A. Panels of such reagents can identify tissue by species and/or by organ type. In a similar fashion, these reagents can be used to screen tissue cultures for contamination. Additional non-limiting examples of such uses are further described herein.

[0359] Because immune/hematopoietic antigens are found expressed in hematopoietic tissues (e.g., bone marrow, fetal liver, and fetal spleen) or cells and tissues of the immune system (e.g., lymph nodes, spleen, B cells, T cells, monocytes, macrophages, dendritic cells, neutrophils, mast cells, basophils, and eosinophils), the polynucleotides of the present invention are also useful as hybridization probes for differential identification of the tissue(s) or cell type(s) present in a biological sample. Similarly, polypeptides and antibodies directed to polypeptides of the present invention are useful to provide immunological probes for differential identification of the tissue(s) (e.g., immunohistochemistry assays) or cell type(s) (e.g., immunocytochemistry assays). In a specific embodiment, the polynucleotides of the present invention are also useful as hybridization probes for differential identification of immune/hematopoietic tissue(s) or cell type(s) present in a biological sample. Similarly, polypeptides and antibodies directed to polypeptides of the present invention are useful to provide immunological probes for differential identification of immune/hematopoietic tissue(s) (e.g., immunohistochemistry assays) or cell type(s) (e.g., immunocytochemistry assays). In addition, for a number of disorders of the above tissues or cells, significantly higher or lower levels of gene expression of the polynucleotides/polypeptides of the present invention may be detected in certain tissues (e.g., tissues expressing polypeptides and/or polynucleotides of the present invention, for example, normal immune/hematopoietic or diseased immune/hematopoietic tissues, and/or those tissues/cells corresponding to the library source relating to a polynucleotide sequence of the invention as disclosed in column 7 of Table 1A, and/or cancerous and/or wounded tissues) or bodily fluids (e.g., semen, lymph, vaginal pool,

serum, plasma, urine, synovial fluid or spinal fluid) taken from an individual having such a disorder, relative to a "standard" gene expression level, i.e., the expression level in healthy tissue from an individual not having the disorder.

[0360] Thus, the invention provides a diagnostic method of a disorder, which involves: (a) assaying gene expression level in cells or body fluid of an individual; (b) comparing the gene expression level with a standard gene expression level, whereby an increase or decrease in the assayed gene expression level compared to the standard expression level is indicative of a disorder.

[0361] In the very least, the polynucleotides of the present invention can be used as molecular weight markers on Southern gels, as diagnostic probes for the presence of a specific mRNA in a particular cell type, as a probe to "subtract-out" known sequences in the process of discovering novel polynucleotides, for selecting and making oligomers for attachment to a "gene chip" or other support, to raise anti-DNA antibodies using DNA immunization techniques, and as an antigen to elicit an immune response.

Uses of the Polypeptides

- [0362] Each of the polypeptides identified herein can be used in numerous ways. The following description should be considered exemplary and utilizes known techniques.
- [0363] Polypeptides and antibodies directed to polypeptides of the present invention are useful to provide immunological probes for differential identification of the tissue(s) (e.g., immunohistochemistry assays such as, for example, ABC immunoperoxidase (Hsu et al., J. Histochem. Cytochem. 29:577-580 (1981)) or cell type(s) (e.g., immunocytochemistry assays).
- [0364] Antibodies can be used to assay levels of polypeptides encoded by polynucleotides of the invention in a biological sample using classical immunohistological methods known to those of skill in the art (see, e.g., Jalkanen, et al., J. Cell. Biol. 101:976-985 (1985); Jalkanen, et al., J. Cell. Biol. 105:3087-3096 (1987)). Other antibody-based methods useful for detecting protein gene expression include immunoassays, such as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). Suitable antibody assay labels are known in the art and include enzyme labels, such as, glucose oxidase; radioisotopes, such as iodine (131 I, 125 I, 123 I, 121 I), carbon (14C), sulfur (35S), tritium (3H), indium (115mIn, 113mIn, 112 In, 111 In), and technetium

(⁹⁹Tc, ^{99m}Tc), thallium (²⁰¹Ti), gallium (⁶⁸Ga, ⁶⁷Ga), palladium (¹⁰³Pd), molybdenum (⁹⁹Mo), xenon (¹³³Xe), fluorine (¹⁸F), ¹⁵³Sm, ¹⁷⁷Lu, ¹⁵⁹Gd, ¹⁴⁹Pm, ¹⁴⁰La, ¹⁷⁵Yb, ¹⁶⁶Ho, ⁹⁰Y, ⁴⁷Sc, ¹⁸⁶Re, ¹⁸⁸Re, ¹⁴²Pr, ¹⁰⁵Rh, ⁹⁷Ru; luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

[0365] In addition to assaying levels of polypeptide of the present invention in a biological sample, proteins can also be detected *in vivo* by imaging. Antibody labels or markers for *in vivo* imaging of protein include those detectable by X-radiography, NMR or ESR. For X-radiography, suitable labels include radioisotopes such as barium or cesium, which emit detectable radiation but are not overtly harmful to the subject. Suitable markers for NMR and ESR include those with a detectable characteristic spin, such as deuterium, which may be incorporated into the antibody by labeling of nutrients for the relevant hybridoma.

[0366] An immune/hematopoietic antigen-specific antibody or antibody fragment which has been labeled with an appropriate detectable imaging moiety, such as a radioisotope (for example, ¹³¹I, ¹¹²In, ^{99m}Tc, (¹³¹I, ¹²⁵I, ¹²³I, ¹²¹I), carbon (¹⁴C), sulfur (³⁵S), tritium (3H), indium (115mIn, 111mIn, 111In), and technetium (99Tc, 99mTc), thallium (201Ti), gallium (68Ga, 67Ga), palladium (103Pd), molybdenum (99Mo), xenon (133Xe), fluorine (¹⁸F, ¹⁵³Sm, ¹⁷⁷Lu, ¹⁵⁹Gd, ¹⁴⁹Pm, ¹⁴⁰La, ¹⁷⁵Yb, ¹⁶⁶Ho, ⁹⁰Y, ⁴⁷Sc, ¹⁸⁶Re, ¹⁸⁸Re, ¹⁴²Pr, 105Rh, 97Ru), a radio-opaque substance, or a material detectable by nuclear magnetic resonance, is introduced (for example, parenterally, subcutaneously or intraperitoneally) into the mammal to be examined for an immune system or hematopoietic disorder. It will be understood in the art that the size of the subject and the imaging system used will determine the quantity of imaging moiety needed to produce diagnostic images. In the case of a radioisotope moiety, for a human subject, the quantity of radioactivity injected will normally range from about 5 to 20 millicuries of 99mTc. The labeled antibody or antibody fragment will then preferentially accumulate at the location of cells which express the polypeptide encoded by a polynucleotide of the invention. In vivo tumor imaging is described in S.W. Burchiel et al., "Immunopharmacokinetics of Radiolabeled Antibodies and Their Fragments" (Chapter 13 in Tumor Imaging: The Radiochemical Detection of Cancer, S.W. Burchiel and B. A. Rhodes, eds., Masson Publishing Inc. (1982)).

In one embodiment, the invention provides a method for the specific delivery of compositions of the invention to cells by administering polypeptides of the invention (e.g., polypeptides encoded by polynucleotides of the invention and/or antibodies) that are associated with heterologous polypeptides or nucleic acids. In one example, the invention provides a method for delivering a therapeutic protein into the targeted cell. In another example, the invention provides a method for delivering a single stranded nucleic acid (e.g., antisense or ribozymes) or double stranded nucleic acid (e.g., DNA that can integrate into the cell's genome or replicate episomally and that can be transcribed) into the targeted cell.

- [0368] In another embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of tumor cells) by administering polypeptides of the invention in association with toxins or cytotoxic prodrugs.
- [0369] In a preferred embodiment, the invention provides a method for the specific destruction of immune/hematopoietic cells (e.g., aberrant immune/hematopoietic cells, immune/hematopoietic neoplasm) by administering polypeptides of the invention (e.g., polypeptides encoded by polynucleotides of the invention and/or antibodies) in association with toxins or cytotoxic prodrugs. In another preferred embodiment the invention provides a method for the specific destruction of tissues/cells corresponding to the library source relating to a polynucleotide sequence of the invention as disclosed in column 7 of Table 1A by administering polypeptides of the invention in association with toxins or cytotoxic prodrugs.
- [0370] By "toxin" is meant one or more compounds that bind and activate endogenous cytotoxic effector systems, radioisotopes, holotoxins, modified toxins, catalytic subunits of toxins, or any molecules or enzymes not normally present in or on the surface of a cell that under defined conditions cause the cell's death. Toxins that may be used according to the methods of the invention include, but are not limited to, radioisotopes known in the art, compounds such as, for example, antibodies (or complement fixing containing portions thereof) that bind an inherent or induced endogenous cytotoxic effector system, thymidine kinase, endonuclease, RNAse, alpha toxin, ricin, abrin, *Pseudomonas* exotoxin A, diphtheria toxin, saporin, momordin, gelonin, pokeweed antiviral protein, alpha-sarcin and cholera toxin. "Toxin" also includes a cytostatic or cytocidal agent, a therapeutic agent or a radioactive metal ion, e.g., alpha-emitters such as, for example, ²¹³Bi, or other

radioisotopes such as, for example, ¹⁰³Pd, ¹³³Xe, ¹³¹I, ¹¹¹In, ⁶⁸Ge, ⁵⁷Co, ⁶⁵Zn, ⁸⁵Sr, ³²P, ³⁵S, ⁹⁰Y, ¹⁵³Sm, ¹⁵³Gd, ¹⁶⁹Yb, ⁵¹Cr, ⁵⁴Mn, ⁷⁵Se, ¹¹³Sn, ⁹⁰Yttrium, ¹¹⁷Tin, ¹⁸⁶Rhenium, ¹⁶⁶Holmium, and ¹⁸⁸Rhenium; luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

- In a specific embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of tumor cells) by administering polypeptides of the invention or antibodies of the invention in association with the radioisotope ⁹⁰Y. In another specific embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of tumor cells) by administering polypeptides of the invention or antibodies of the invention in association with the radioisotope ¹¹¹In. In a further specific embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of tumor cells) by administering polypeptides of the invention or antibodies of the invention in association with the radioisotope ¹³¹I.
- [0372] Techniques known in the art may be applied to label polypeptides of the invention (including antibodies). Such techniques include, but are not limited to, the use of bifunctional conjugating agents (see e.g., U.S. Patent Nos. 5,756,065; 5,714,631; 5,696,239; 5,652,361; 5,505,931; 5,489,425; 5,435,990; 5,428,139; 5,342,604; 5,274,119; 4,994,560; and 5,808,003; the contents of each of which are hereby incorporated by reference in its entirety).
- [0373] Thus, the invention provides a diagnostic method of a disorder, which involves (a) assaying the expression level of a polypeptide of the present invention in cells or body fluid of an individual; and (b) comparing the assayed polypeptide expression level with a standard polypeptide expression level, whereby an increase or decrease in the assayed polypeptide expression level compared to the standard expression level is indicative of a disorder. With respect to cancer, the presence of a relatively high amount of transcript in biopsied tissue from an individual may indicate a predisposition for the development of the disease, or may provide a means for detecting the disease prior to the appearance of actual clinical symptoms. A more definitive diagnosis of this type may allow health professionals to employ preventative measures or aggressive treatment earlier thereby preventing the development or further progression of the cancer.
- [0374] Moreover, polypeptides of the present invention can be used to treat or prevent diseases or conditions of/assocatied with hematopoiesis and/or the immune system such

as, for example, such as anemias, autoimmune diseases, immunodeficiencies, allergies, asthma, and inflammation) and infectious/parasitic diseases. In preferred embodiments, polynucleotides expressed in a particular tissue type (see, e.g., Table 1A, column 7) are used to diagnose, detect, prevent, treat and/or prognose disorders associated with the tissue type. For example, patients can be administered a polypeptide of the present invention in an effort to replace absent or decreased levels of the polypeptide (e.g., insulin), to supplement absent or decreased levels of a different polypeptide (e.g., hemoglobin S for hemoglobin B, SOD, catalase, DNA repair proteins), to inhibit the activity of a polypeptide (e.g., an oncogene or tumor supressor), to activate the activity of a polypeptide (e.g., by binding to a receptor), to reduce the activity of a membrane bound receptor by competing with it for free ligand (e.g., soluble TNF receptors used in reducing inflammation), or to bring about a desired response (e.g., blood vessel growth inhibition, enhancement of the immune response to proliferative cells or tissues).

[0375] Similarly, antibodies directed to a polypeptide of the present invention can also be used to treat disease (as described *supra*, and elsewhere herein). For example, administration of an antibody directed to a polypeptide of the present invention can bind, and/or neutralize the polypeptide, and/or reduce overproduction of the polypeptide. Similarly, administration of an antibody can activate the polypeptide, such as by binding to a polypeptide bound to a membrane (receptor).

[0376] At the very least, the polypeptides of the present invention can be used as molecular weight markers on SDS-PAGE gels or on molecular sieve gel filtration columns using methods well known to those of skill in the art. Polypeptides can also be used to raise antibodies, which in turn are used to measure protein expression from a recombinant cell, as a way of assessing transformation of the host cell. Moreover, the polypeptides of the present invention can be used to test the biological activities described herein.

Diagnostic Asssays

[0377] The compounds of the present invention are useful for diagnosis, treatment, prevention and/or prognosis of various immune/hematopoietic related disorders in mammals, preferably humans. Such disorders include, but are not limited to such as anemias, autoimmune diseases, immunodeficiencies, allergies, asthma, and inflammation) and infectious/parasitic diseases. In preferred embodiments, polynucleotides expressed in

a particular tissue type (see, e.g., Table 1A, column 7) are used to diagnose, detect, prevent, treat and/or prognose disorders associated with the tissue type.

[0378]Immune/hematopoietic antigens are expressed hematopoietic tissues (e.g., bone marrow, fetal liver, and fetal spleen) or cells and tissues of the immune system (e.g., lymph nodes, spleen, B cells, T cells, monocytes, macrophages, dendritic cells, neutrophils, mast cells, basophils, and eosinophils). For a number of immune/hematopoietic-related disorders, substantially altered (increased or decreased) levels of immune/hematopoietic antigen gene expression can be detected in immune/hematopoietic tissue or other cells or bodily fluids (e.g., sera, plasma, urine, semen, synovial fluid or spinal fluid) taken from an individual having such a disorder, relative to a "standard" immune/hematopoietic antigen gene expression level, that is, the immune/hematopoietic antigen expression level in immune/hematopoietic tissues or bodily fluids from an individual not having the immune/hematopoietic-related disorder. Thus, the invention provides a diagnostic method useful during diagnosis of an immune/hematopoietic-related disorder, which involves measuring the expression level of the gene encoding the immune/hematopoietic associated polypeptide in immune/hematopoietic tissue or other cells or body fluid from an individual and comparing the measured gene expression level with a standard immune/hematopoietic antigens gene expression level, whereby an increase or decrease in the gene expression level(s) compared to the standard is indicative of an immune/hematopoietic-related disorder.

In specific embodiments, the invention provides a diagnostic method useful during diagnosis of a disorder of a normal or diseased tissue/cell source corresponding to column 7 of Table 1A, which involves measuring the expression level of the coding sequence of a polynucleotide sequence associated with this tissue/cell source as disclosed in Table 1A in the tissue/cell source or other cells or body fluid from an individual and comparing the expression level of the coding sequence with a standard expression level of the coding sequence of a polynucleotide sequence, whereby an increase or decrease in the gene expression level(s) compared to the standard is indicative of a disorder of a normal or diseased tissue/cell source corresponding to column 7 of Table 1A.

[0380] In particular, it is believed that certain tissues in mammals with cancer of cells of hematopoietic origin or cancer of tissues of the immune system (e.g., lymph nodes, spleen.

bone marrow) express significantly enhanced or reduced levels of normal or altered immune/hematopoietic antigen expression and mRNA encoding the immune/hematopoietic associated polypeptide when compared to a corresponding "standard" level. Further, it is believed that enhanced or depressed levels of the immune/hematopoietic associated polypeptide can be detected in certain body fluids (e.g., sera, plasma, urine, and spinal fluid) or cells or tissue from mammals with such a cancer when compared to sera from mammals of the same species not having the cancer.

For example, as disclosed herein, immune/hematopoietic associated [0381] polypeptides of the invention are expressed in hematopoietic tissues (e.g., bone marrow, fetal liver, and fetal spleen) or cells and tissues of the immune system (e.g., lymph nodes, spleen, B cells, T cells, monocytes, macrophages, dendritic cells, neutrophils, mast cells, basophils, and eosinophils). Accordingly, polynucleotides of the invention (e.g., polynucleotide sequences complementary to all or a portion of an immune/hematopoietic antigen mRNA nucleotide sequence of SEQ ID NO:X, nucleotide sequence encoding SEQ ID NO:Y, nucleotide sequence encoding a polypeptide encoded by SEQ ID NO:X and/or a nucleotide sequence delineated by columns 8 and 9 of Table 2) and antibodies (and antibody fragments) directed against the polypeptides of the invention may be used to quantitate or qualitate concentrations of cells expressing immune/hematopoietic antigens, preferably on their cell surfaces. These polynucleotides and antibodies additionally have diagnostic applications in detecting abnormalities in the level of immune/hematopoietic antigens gene expression, or abnormalities in the structure and/or temporal, tissue, cellular, or subcellular location of immune/hematopoietic antigens. These diagnostic assays may be performed in vivo or in vitro, such as, for example, on blood samples, biopsy tissue or autopsy tissue. In specific embodiments, polynucleotides and antibodies of the invention are used to quantitate or qualitate tissues/cells corresponding to the library source disclosed in column 7 of Table 1A expressing the corresponding immune/hematopoietic sequence disclosed in the same row of Table 1A, preferrably on their cell surface.

[0382] Thus, the invention provides a diagnostic method useful during diagnosis of an immune/hematopoietic-related disorder, including cancers, which involves measuring the expression level of the gene encoding the immune/hematopoietic antigen polypeptide in immune/hematopoietic tissue or other cells or body fluid from an individual and

comparing the measured gene expression level with a standard immune/hematopoietic antigen gene expression level, whereby an increase or decrease in the gene expression level compared to the standard is indicative of an immune/hematopoietic-related disorder. In specific embodiments, polynucleotides and antibodies of the invention are used to quantitate or qualitate tissues/cells corresponding to the library source disclosed in column 7 of Table 1A expressing the corresponding immune/hematopoietic sequence disclosed in the same row of Table 1A, preferrably on their cell surface.

[0383] Where a diagnosis of an immune/hematopoietic-related disease or disorder, including diagnosis of a tumor, has already been made according to conventional methods, the present invention is useful as a prognostic indicator, whereby patients exhibiting enhanced or depressed immune/hematopoietic antigen gene expression will experience a worse clinical outcome relative to patients expressing the gene at a level nearer the standard level.

By "assaying the expression level of the gene encoding the [0384] immune/hematopoietic associated polypeptide" is intended qualitatively or quantitatively measuring or estimating the level of the immune/hematopoietic antigen polypeptide or the level of the mRNA encoding the immune/hematopoietic antigen polypeptide in a first biological sample either directly (e.g., by determining or estimating absolute protein level or mRNA level) or relatively (e.g., by comparing to the immune/hematopoietic associated polypeptide level or mRNA level in a second biological sample). Preferably, the immune/hematopoietic antigen polypeptide expression level or mRNA level in the first biological sample is measured or estimated and compared to a standard immune/hematopoietic antigen polypeptide level or mRNA level, the standard being taken from a second biological sample obtained from an individual not having the disorder or being determined by averaging levels from a population of individuals not having an immune/hematopoietic -related disorder. As will be appreciated in the art, once a standard immune/hematopoietic antigen polypeptide level or mRNA level is known, it can be used repeatedly as a standard for comparison.

[0385] By "biological sample" is intended any biological sample obtained from an individual, cell line, tissue culture, or other source containing immune/hematopoietic antigen polypeptides (including portions thereof) or mRNA. As indicated, biological samples include body fluids (such as sera, plasma, urine, synovial fluid and spinal fluid)

which contain cells expressing immune/hematopoietic antigen polypeptides, heamtopoietic cells and tissue, cells and tissues of the immune system, and other tissue sources found to express the full length or fragments thereof of an immune/hematopoietic antigen. Methods for obtaining tissue biopsies and body fluids from mammals are well known in the art. Where the biological sample is to include mRNA, a tissue biopsy is the preferred source.

[0386] Total cellular RNA can be isolated from a biological sample using any suitable technique such as the single-step guanidinium-thiocyanate-phenol-chloroform method described in Chomczynski and Sacchi, Anal. Biochem. 162:156-159 (1987). Levels of mRNA encoding the immune/hematopoietic antigen polypeptides are then assayed using any appropriate method. These include Northern blot analysis, S1 nuclease mapping, the polymerase chain reaction (PCR), reverse transcription in combination with the polymerase chain reaction (RT-PCR), and reverse transcription in combination with the ligase chain reaction (RT-LCR).

[0387] The present invention also relates to diagnostic assays such as quantitative and diagnostic assays for detecting levels of immune/hematopoietic antigen polypeptides, in a biological sample (e.g., cells and tissues), including determination of normal and abnormal levels of polypeptides. Thus, for instance, a diagnostic assay in accordance with the invention for detecting over-expression of immune/hematopoietic antigens compared to normal control tissue samples may be used to detect the presence of tumors. Assay techniques that can be used to determine levels of a polypeptide, such as an immune/hematopoietic antigen polypeptide of the present invention in a sample derived from a host are well-known to those of skill in the art. Such assay methods include radioimmunoassays, competitive-binding assays, Western Blot analysis and ELISA assays. Assaying immune/hematopoietic antigen polypeptide levels in a biological sample can occur using any art-known method.

[0388] Assaying immune/hematopoietic antigen polypeptide levels in a biological sample can occur using antibody-based techniques. For example, immune/hematopoietic antigen polypeptide expression in tissues can be studied with classical immunohistological methods (Jalkanen et al., J. Cell. Biol. 101:976-985 (1985); Jalkanen, M., et al., J. Cell. Biol. 105:3087-3096 (1987)). Other antibody-based methods useful for detecting immune/hematopoietic antigen polypeptide gene expression include immunoassays, such

as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). Suitable antibody assay labels are known in the art and include enzyme labels, such as, glucose oxidase, and radioisotopes, such as iodine (¹²⁵I, ¹²¹I), carbon (¹⁴C), sulfur (³⁵S), tritium (³H), indium (¹¹²In), and technetium (^{99m}Tc), and fluorescent labels, such as fluorescein and rhodamine, and biotin.

- [0389] The tissue or cell type to be analyzed will generally include those which are known, or suspected, to express the immune/hematopoietic antigen gene (such as, for example, cells of the immune system or cancers of cells of hematopoetic origin (e.g., leukemias). The protein isolation methods employed herein may, for example, be such as those described in Harlow and Lane (Harlow, E. and Lane, D., 1988, "Antibodies: A Laboratory Manual", Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York), which is incorporated herein by reference in its entirety. The isolated cells can be derived from cell culture or from a patient. The analysis of cells taken from culture may be a necessary step in the assessment of cells that could be used as part of a cell-based gene therapy technique or, alternatively, to test the effect of compounds on the expression of the immune/hematopoietic antigen gene.
- [0390] For example, antibodies, or fragments of antibodies, such as those described herein, may be used to quantitatively or qualitatively detect the presence of immune/hematopoietic antigen gene products or conserved variants or peptide fragments thereof. This can be accomplished, for example, by immunofluorescence techniques employing a fluorescently labeled antibody coupled with light microscopic, flow cytometric, or fluorimetric detection.
- In a preferred embodiment, antibodies, or fragments of antibodies directed to any one or all of the predicted epitope domains of the immune/hematopoietic antigen polypeptides (Shown in Table 1A, column 6) may be used to quantitatively or qualitatively detect the presence of immune/hematopoietic antigen gene products or conserved variants or peptide fragments thereof. This can be accomplished, for example, by immunofluorescence techniques employing a fluorescently labeled antibody coupled with light microscopic, flow cytometric, or fluorimetric detection.
- [0392] In an additional preferred embodiment, antibodies, or fragments of antibodies directed to a conformational epitope of an immune/hematopoietic antigen may be used to quantitatively or qualitatively detect the presence of immune/hematopoietic antigen gene

products or conserved variants or peptide fragments thereof. This can be accomplished, for example, by immunofluorescence techniques employing a fluorescently labeled antibody coupled with light microscopic, flow cytometric, or fluorimetric detection.

[0393] The antibodies (or fragments thereof), and/or immune/hematopoietic antigen polypeptides of the present invention may, additionally, be employed histologically, as in immunofluorescence, immunoelectron microscopy or non-immunological assays, for in situ detection of immune/hematopoietic antigen gene products or conserved variants or peptide fragments thereof. In situ detection may be accomplished by removing a histological specimen from a patient, and applying thereto a labeled antibody or immune/hematopoietic antigen polypeptide of the present invention. The antibody (or fragment thereof) or immune/hematopoietic antigen polypeptide is preferably applied by overlaying the labeled antibody (or fragment) onto a biological sample. Through the use of such a procedure, it is possible to determine not only the presence of the immune/hematopoietic antigen gene product, or conserved variants or peptide fragments. or immune/hematopoietic antigen polypeptide binding, but also its distribution in the examined tissue. Using the present invention, those of ordinary skill will readily perceive that any of a wide variety of histological methods (such as staining procedures) can be modified in order to achieve such in situ detection.

[0394] Immunoassays and non-immunoassays for immune/hematopoietic antigen gene products or conserved variants or peptide fragments thereof will typically comprise incubating a sample, such as a biological fluid, a tissue extract, freshly harvested cells, or lysates of cells which have been incubated in cell culture, in the presence of a detectably labeled antibody capable of binding immune/hematopoietic antigen gene products or conserved variants or peptide fragments thereof, and detecting the bound antibody by any of a number of techniques well-known in the art.

[0395] The biological sample may be brought in contact with and immobilized onto a solid phase support or carrier such as nitrocellulose, or other solid support which is capable of immobilizing cells, cell particles or soluble proteins. The support may then be washed with suitable buffers followed by treatment with the detectably labeled anti-immune/hematopoietic antigen antibody or detectable immune/hematopoietic antigen polypeptide. The solid phase support may then be washed with the buffer a second time to remove unbound antibody or polypeptide. Optionally the antibody is subsequently

labeled. The amount of bound label on solid support may then be detected by conventional means.

[0396] By "solid phase support or carrier" is intended any support capable of binding an antigen or an antibody. Well-known supports or carriers include glass, polystyrene, polypropylene, polyethylene, dextran, nylon, amylases, natural and modified celluloses, polyacrylamides, gabbros, and magnetite. The nature of the carrier can be either soluble to some extent or insoluble for the purposes of the present invention. The support material may have virtually any possible structural configuration so long as the coupled molecule is capable of binding to an antigen or antibody. Thus, the support configuration may be spherical, as in a bead, or cylindrical, as in the inside surface of a test tube, or the external surface of a rod. Alternatively, the surface may be flat such as a sheet, test strip, etc. Preferred supports include polystyrene beads. Those skilled in the art will know many other suitable carriers for binding antibody or antigen, or will be able to ascertain the same by use of routine experimentation.

[0397] The binding activity of a given lot of anti- immune/hematopoietic antigen antibody or immune/hematopoietic antigen polypeptide may be determined according to well known methods. Those skilled in the art will be able to determine operative and optimal assay conditions for each determination by employing routine experimentation.

In addition to assaying immune/hematopoietic antigen polypeptide levels or polynucleotide levels in a biological sample obtained from an individual, immune/hematopoietic antigen polypeptide or polynucleotide can also be detected *in vivo* by imaging. For example, in one embodiment of the invention, immune/hematopoietic antigen polypeptide and/or anti-immune/hematopoietic antigen antibodies are used to image immune system diseased cells or diseased hematopoietic cells, such as neoplasms. In another embodiment, immune/hematopoietic antigen polynucleotides of the invention (e.g., polynucleotides complementary to all or a portion of immune/hematopoietic antigen mRNA) and/or anti-immune/hematopoietic antigen antibodies (e.g., antibodies directed to any one or a combination of the epitopes of immune/hematopoietic antigens, antibodies directed to a conformational epitope of immune/hematopoietic antigens, antibodies directed to the full length polypeptide expressed on the cell surface of a mammalian cell) are used to image diseased or neoplastic cells of the immune system or disease or neoplastic cells assoctiated with hematopoiesis.

[0399] Antibody labels or markers for in vivo imaging of immune/hematopoietic antigen polypeptides include those detectable by X-radiography, NMR, MRI, CAT-scans or ESR. For X-radiography, suitable labels include radioisotopes such as barium or cesium, which emit detectable radiation but are not overtly harmful to the subject. Suitable markers for NMR and ESR include those with a detectable characteristic spin, such as deuterium, which may be incorporated into the antibody by labeling of nutrients for the relevant hybridoma. Where in vivo imaging is used to detect enhanced levels of immune/hematopoietic antigen polypeptides for diagnosis in humans, it may be preferable to use human antibodies or "humanized" chimeric monoclonal antibodies. antibodies can be produced using techniques described herein or otherwise known in the art. For example methods for producing chimeric antibodies are known in the art. See, for review, Morrison, Science 229:1202 (1985); Oi et al., BioTechniques 4:214 (1986); Cabilly et al., U.S. Patent No. 4,816,567; Taniguchi et al., EP 171496; Morrison et al., EP 173494; Neuberger et al., WO 8601533; Robinson et al., WO 8702671; Boulianne et al., Nature 312:643 (1984); Neuberger et al., Nature 314:268 (1985).

[0400] Additionally, any immune/hematopoietic antigen polypeptides whose presence can be detected, can be administered. For example, immune/hematopoietic antigen polypeptides labeled with a radio-opaque or other appropriate compound can be administered and visualized *in vivo*, as discussed, above for labeled antibodies. Further such immune/hematopoietic antigen polypeptides can be utilized for *in vitro* diagnostic procedures.

fragment which has been labeled with an appropriate detectable imaging moiety, such as a radioisotope (for example, ¹³¹I, ¹¹²In, ^{99m}Tc), a radio-opaque substance, or a material detectable by nuclear magnetic resonance, is introduced (for example, parenterally, subcutaneously or intraperitoneally) into the mammal to be examined for an immune/hematopoietic-related disorder. It will be understood in the art that the size of the subject and the imaging system used will determine the quantity of imaging moiety needed to produce diagnostic images. In the case of a radioisotope moiety, for a human subject, the quantity of radioactivity injected will normally range from about 5 to 20 millicuries of ^{99m}Tc. The labeled antibody or antibody fragment will then preferentially accumulate at the location of cells which contain immune/hematopoietic antigen protein.

In vivo tumor imaging is described in S.W. Burchiel et al., "Immunopharmacokinetics of Radiolabeled Antibodies and Their Fragments" (Chapter 13 in *Tumor Imaging: The Radiochemical Detection of Cancer*, S.W. Burchiel and B. A. Rhodes, eds., Masson Publishing Inc. (1982)).

[0402]With respect to antibodies, one of the ways in which the antiimmune/hematopoietic antigen antibody can be detectably labeled is by linking the same to an enzyme and using the linked product in an enzyme immunoassay (EIA) (Voller, A., "The Enzyme Linked Immunosorbent Assay (ELISA)", 1978, Diagnostic Horizons 2:1-7. Microbiological Associates Quarterly Publication, Walkersville, MD); Voller et al., J. Clin. Pathol. 31:507-520 (1978); Butler, J.E., Meth. Enzymol. 73:482-523 (1981); Maggio, E. (ed.), 1980, Enzyme Immunoassay, CRC Press, Boca Raton, FL,; Ishikawa, E. et al., (eds.), 1981, Enzyme Immunoassay, Kgaku Shoin, Tokyo). The enzyme which is bound to the antibody will react with an appropriate substrate, preferably a chromogenic substrate, in such a manner as to produce a chemical moiety which can be detected, for example, by spectrophotometric, fluorimetric or by visual means. Enzymes which can be used to detectably label the antibody include, but are not limited to, malate dehydrogenase, staphylococcal nuclease, delta-5-steroid isomerase, yeast alcohol dehydrogenase, alpha-glycerophosphate, dehydrogenase, triose phosphate isomerase, horseradish peroxidase, alkaline phosphatase, asparaginase, glucose oxidase, betagalactosidase, ribonuclease, urease, catalase, glucose-6-phosphate dehydrogenase, glucoamylase and acetylcholinesterase. Additionally, the detection can be accomplished by colorimetric methods which employ a chromogenic substrate for the enzyme. Detection may also be accomplished by visual comparison of the extent of enzymatic reaction of a substrate in comparison with similarly prepared standards.

[0403] Detection may also be accomplished using any of a variety of other immunoassays. For example, by radioactively labeling the antibodies or antibody fragments, it is possible to detect immune/hematopoietic antigens through the use of a radioimmunoassay (RIA) (see, for example, Weintraub, B., Principles of Radioimmunoassays, Seventh Training Course on Radioligand Assay Techniques, The Endocrine Society, March, 1986, which is incorporated by reference herein). The radioactive isotope can be detected by means including, but not limited to, a gamma counter, a scintillation counter, or autoradiography.

[0404] It is also possible to label the antibody with a fluorescent compound. When the fluorescently labeled antibody is exposed to light of the proper wave length, its presence can then be detected due to fluorescence. Among the most commonly used fluorescent labeling compounds are fluorescein isothiocyanate, rhodamine, phycocythrin, phycocyanin, allophycocyanin, ophthaldehyde and fluorescemine.

- [0405] The antibody can also be detectably labeled using fluorescence emitting metals such as ¹⁵²Eu, or others of the lanthanide series. These metals can be attached to the antibody using such metal chelating groups as diethylenetriaminepentacetic acid (DTPA) or ethylenediaminetetraacetic acid (EDTA).
- [0406] The antibody also can be detectably labeled by coupling it to a chemiluminescent compound. The presence of the chemiluminescent-tagged antibody is then determined by detecting the presence of luminescence that arises during the course of a chemical reaction. Examples of particularly useful chemiluminescent labeling compounds are luminol, isoluminol, theromatic acridinium ester, imidazole, acridinium salt and oxalate ester.
- [0407] Likewise, a bioluminescent compound may be used to label the antibody of the present invention. Bioluminescence is a type of chemiluminescence found in biological systems in, which a catalytic protein increases the efficiency of the chemiluminescent reaction. The presence of a bioluminescent protein is determined by detecting the presence of luminescence. Important bioluminescent compounds for purposes of labeling are luciferin, luciferase and aequorin.

Methods for Detecting Immune/Hematopoietic-Related Disease, Including Cancer

In general, an immune/hematopoietic-related disease or cancer may be detected in a patient based on the presence of one or more immune/hematopoietic antigen proteins of the invention and/or polynucleotides encoding such proteins in a biological sample (for example, blood, sera, urine, and/or tumor biopsies) obtained from the patient. In other words, such proteins and/or polynucleotides may be used as markers to indicate the presence or absence of an immune/hematopoietic-related disease or disorder, including cancer. Cancers that may be diagnosed, and/or prognosed using the compositions of the invention include but are not limited to, cancers of cells of hematopoietic origins such as leukemias and lymphomas. In addition, such proteins and/or polynucleotids may be useful

for the detection of other diseases and cancers, including cancers of tissues/cells corresponding to the library source disclosed in column 7 of Table 1A expressing the corresponding immune/hematopoietic sequence disclosed in the same row of Table 1A. The binding agents provided herein generally permit detection of the level of antigen that binds to the agent in the biological sample. Polynucleotide primers and probes may be used to detect the level of mRNA encoding immune/hematopoietic antigen polypeptides, which is also indicative of the presence or absence of an immune/hematopoietic-related disease or disorder, including cancer. In general, immune/hematopoietic antigen polypeptides should be present at a level that is at least three fold higher in diseased tissue than in normal tissue.

There are a variety of assay formats known to those of ordinary skill in the art for using a binding agent to detect polypeptide markers in a sample. See, e.g., Harlow and Lane, supra. In general, the presence or absence of an immune/hematopoietic-related disease in a patient may be determined by (a) contacting a biological sample obtained from a patient with a binding agent; (b) detecting in the sample a level of polypeptide that binds to the binding agent; and (c) comparing the level of polypeptide with a predetermined cut-off value.

[0410] In a preferred embodiment, the assay involves the use of binding agent immobilized on a solid support to bind to and remove the immune/hematopoietic antigen polypeptide of the invention from the remainder of the sample. The bound polypeptide may then be detected using a detection reagent that contains a reporter group and specifically binds to the binding agent/polypeptide complex. Such detection reagents may comprise, for example, a binding agent that specifically binds to the polypeptide or an antibody or other agent that specifically binds to the binding agent, such as an antiimmunoglobulin, protein G, protein A or a lectin. Alternatively, a competitive assay may be utilized, in which a polypeptide is labeled with a reporter group and allowed to bind to the immobilized binding agent after incubation of the binding agent with the sample. The extent to which components of the sample inhibit the binding of the labeled polypeptide to the binding agent is indicative of the reactivity of the sample with the immobilized binding agent. Suitable polypeptides for use within such assays include immune/hematopoietic antigen polypeptides and portions thereof, or antibodies, to which the binding agent binds, as described above.

The solid support may be any material known to those of skill in the art to [0411] which immune/hematopoietic antigen polypeptides of the invention may be attached. For example, the solid support may be a test well in a microtiter plate or a nitrocellulose or other suitable membrane. Alternatively, the support may be a bead or disc, such as glass fiberglass, latex or a plastic material such as polystyrene or polyvinylchloride. The support may also be a magnetic particle or a fiber optic sensor, such as those disclosed, for example, in U.S. Patent No. 5,359,681. The binding agent may be immobilized on the solid support using a variety of techniques known to those of skill in the art, which are amply described in the patent and scientific literature. In the context of the present invention, the term "immobilization" refers to both noncovalent association, such as adsorption, and covalent attachment (which may be a direct linkage between the agent and functional groups on the support or may be a linkage by way of a cross-linking agent). Immobilization by adsorption to a well in a microtiter plate or to a membrane is preferred. In such cases, adsorption may be achieved by contacting the binding agent, in a suitable buffer, with the solid support for the suitable amount of time. The contact time varies with temperature, but is typically between about 1 hour and about 1 day. In general, contacting a well of plastic microtiter plate (such as polystyrene or polyvinylchloride) with an amount of binding agent ranging from about 10 ng to about 10 ug, and preferably about 100 ng to about 1 ug, is sufficient to immobilize an adequate amount of binding agent.

[0412] Covalent attachment of binding agent to a solid support may generally be achieved by first reacting the support with a bifunctional reagent that will react with both the support and a functional group, such as a hydroxyl or amino group, on the binding agent. For example, the binding agent may be covalently attached to supports having an appropriate polymer coating using benzoquinone or by condensation of an aldehyde group on the support with an amine and an active hydrogen on the binding partner (see, e.g., Pierce Immunotechnology Catalog and Handbook, 1991, at A12-A13).

Gene Therapy Methods

[0413] Also encompassed by the present invention are gene therapy methods for treating or preventing disorders, diseases and conditions. The gene therapy methods relate to the introduction of nucleic acid (DNA, RNA and antisense DNA or RNA) sequences into an animal to achieve expression of an immune/hematopoietic antigen of the present

invention. This method requires a polynucleotide which codes for a polypeptide of the present invention operatively linked to a promoter and any other genetic elements necessary for the expression of the polypeptide by the target tissue. Such gene therapy and delivery techniques are known in the art, see, for example, WO90/11092, which is herein incorporated by reference.

- (DNA or RNA) comprising a promoter operably linked to a polynucleotide of the present invention ex vivo, with the engineered cells then being provided to a patient to be treated with the polypeptide of the present invention. Such methods are well-known in the art. For example, see Belldegrun, A., et al., J. Natl. Cancer Inst. 85: 207-216 (1993); Ferrantini, M. et al., Cancer Research 53: 1107-1112 (1993); Ferrantini, M. et al., J. Immunology 153: 4604-4615 (1994); Kaido, T., et al., Int. J. Cancer 60: 221-229 (1995); Ogura, H., et al., Cancer Research 50: 5102-5106 (1990); Santodonato, L., et al., Human Gene Therapy 7:1-10 (1996); Santodonato, L., et al., Gene Therapy 4:1246-1255 (1997); and Zhang, J.-F. et al., Cancer Gene Therapy 3: 31-38 (1996)), which are herein incorporated by reference. In one embodiment, the cells which are engineered are arterial cells. The arterial cells may be reintroduced into the patient through direct injection to the artery, the tissues surrounding the artery, or through catheter injection.
- [0415] As discussed in more detail below, the polynucleotide constructs can be delivered by any method that delivers injectable materials to the cells of an animal, such as, injection into the interstitial space of tissues (heart, muscle, skin, lung, liver, and the like). The polynucleotide constructs may be delivered in a pharmaceutically acceptable liquid or aqueous carrier.
- [0416] In one embodiment, the polynucleotide of the present invention is delivered as a naked polynucleotide. The term "naked" polynucleotide, DNA or RNA refers to sequences that are free from any delivery vehicle that acts to assist, promote or facilitate entry into the cell, including viral sequences, viral particles, liposome formulations, lipofectin or precipitating agents and the like. However, the polynucleotide of the present invention can also be delivered in liposome formulations and lipofectin formulations and the like can be prepared by methods well known to those skilled in the art. Such methods are described, for example, in U.S. Patent Nos. 5,593,972, 5,589,466, and 5,580,859, which are herein incorporated by reference.

[0417] The polynucleotide vector constructs used in the gene therapy method are preferably constructs that will not integrate into the host genome nor will they contain sequences that allow for replication. Appropriate vectors include pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; pSVK3, pBPV, pMSG and pSVL available from Pharmacia; and pEF1/V5, pcDNA3.1, and pRc/CMV2 available from Invitrogen. Other suitable vectors will be readily apparent to the skilled artisan.

- [0418] Any strong promoter known to those skilled in the art can be used for driving the expression of the polynucleotide sequence. Suitable promoters include adenoviral promoters, such as the adenoviral major late promoter; or heterologous promoters, such as the cytomegalovirus (CMV) promoter; the respiratory syncytial virus (RSV) promoter; inducible promoters, such as the MMT promoter, the metallothionein promoter; heat shock promoters; the albumin promoter; the ApoAI promoter; human globin promoters; viral thymidine kinase promoters, such as the Herpes Simplex thymidine kinase promoter; retroviral LTRs; the b-actin promoter; and human growth hormone promoters. The promoter also may be the native promoter for the polynucleotide of the present invention.
- [0419] Unlike other gene therapy techniques, one major advantage of introducing naked nucleic acid sequences into target cells is the transitory nature of the polynucleotide synthesis in the cells. Studies have shown that non-replicating DNA sequences can be introduced into cells to provide production of the desired polypeptide for periods of up to six months.
- [0420] The polynucleotide construct can be delivered to the interstitial space of tissues within the an animal, including of muscle, skin, brain, lung, liver, spleen, bone marrow, thymus, heart, lymph, blood, bone, cartilage, pancreas, kidney, gall bladder, stomach, intestine, testis, ovary, uterus, rectum, nervous system, eye, gland, and connective tissue. Interstitial space of the tissues comprises the intercellular, fluid, mucopolysaccharide matrix among the reticular fibers of organ tissues, elastic fibers in the walls of vessels or chambers, collagen fibers of fibrous tissues, or that same matrix within connective tissue ensheathing muscle cells or in the lacunae of bone. It is similarly the space occupied by the plasma of the circulation and the lymph fluid of the lymphatic channels. Delivery to the interstitial space of muscle tissue is preferred for the reasons discussed below. They may be conveniently delivered by injection into the tissues comprising these cells. They are preferably delivered to and expressed in persistent, non-dividing cells which are

differentiated, although delivery and expression may be achieved in non-differentiated or less completely differentiated cells, such as, for example, stem cells of blood or skin fibroblasts. *In vivo* muscle cells are particularly competent in their ability to take up and express polynucleotides.

- [0421] For the naked nucleic acid sequence injection, an effective dosage amount of DNA or RNA will be in the range of from about 0.05 mg/kg body weight to about 50 mg/kg body weight. Preferably the dosage will be from about 0.005 mg/kg to about 20 mg/kg and more preferably from about 0.05 mg/kg to about 5 mg/kg. Of course, as the artisan of ordinary skill will appreciate, this dosage will vary according to the tissue site of injection. The appropriate and effective dosage of nucleic acid sequence can readily be determined by those of ordinary skill in the art and may depend on the condition being treated and the route of administration.
- [0422] The preferred route of administration is by the parenteral route of injection into the interstitial space of tissues. However, other parenteral routes may also be used, such as, inhalation of an aerosol formulation particularly for delivery to lungs or bronchial tissues, throat or mucous membranes of the nose. In addition, naked DNA constructs can be delivered to arteries during angioplasty by the catheter used in the procedure.
- [0423] The naked polynucleotides are delivered by any method known in the art, including, but not limited to, direct needle injection at the delivery site, intravenous injection, topical administration, catheter infusion, and so-called "gene guns". These delivery methods are known in the art.
- [0424] The constructs may also be delivered with delivery vehicles such as viral sequences, viral particles, liposome formulations, lipofectin, precipitating agents, etc. Such methods of delivery are known in the art.
- [0425] In certain embodiments, the polynucleotide constructs are complexed in a liposome preparation. Liposomal preparations for use in the instant invention include cationic (positively charged), anionic (negatively charged) and neutral preparations. However, cationic liposomes are particularly preferred because a tight charge complex can be formed between the cationic liposome and the polyanionic nucleic acid. Cationic liposomes have been shown to mediate intracellular delivery of plasmid DNA (Felgner et al., Proc. Natl. Acad. Sci. USA (1987) 84:7413-7416, which is herein incorporated by reference); mRNA (Malone et al., Proc. Natl. Acad. Sci. USA (1989) 86:6077-6081,

which is herein incorporated by reference); and purified transcription factors (Debs et al., J. Biol. Chem. (1990) 265:10189-10192, which is herein incorporated by reference), in functional form.

- [0426] Cationic liposomes are readily available. For example, N[1-2,3-dioleyloxy)propyl]-N,N,N-triethylammonium (DOTMA) liposomes are particularly useful and are available under the trademark Lipofectin, from GIBCO BRL, Grand Island, N.Y., (see, also, Felgner et al., Proc. Natl Acad. Sci. USA (1987) 84:7413-7416, which is herein incorporated by reference). Other commercially available liposomes include transfectace (DDAB/DOPE) and DOTAP/DOPE (Boehringer).
- [0427] Other cationic liposomes can be prepared from readily available materials using techniques well known in the art. See, e.g. PCT Publication No. WO 90/11092 (which is herein incorporated by reference) for a description of the synthesis of DOTAP (1,2-bis(oleoyloxy)-3-(trimethylammonio)propane) liposomes. Preparation of DOTMA liposomes is explained in the literature, see, e.g., P. Felgner et al., Proc. Natl. Acad. Sci. USA 84:7413-7417, which is herein incorporated by reference. Similar methods can be used to prepare liposomes from other cationic lipid materials.
- [0428] Similarly, anionic and neutral liposomes are readily available, such as from Avanti Polar Lipids (Birmingham, Ala.), or can be easily prepared using readily available materials. Such materials include phosphatidyl choline, cholesterol, phosphatidyl ethanolamine, dioleoylphosphatidyl choline (DOPC), dioleoylphosphatidyl glycerol (DOPG), dioleoylphoshatidyl ethanolamine (DOPE), among others. These materials can also be mixed with the DOTMA and DOTAP starting materials in appropriate ratios. Methods for making liposomes using these materials are well known in the art.
- [0429] For example, commercially dioleoylphosphatidyl choline (DOPC), dioleoylphosphatidyl glycerol (DOPG), and dioleoylphosphatidyl ethanolamine (DOPE) can be used in various combinations to make conventional liposomes, with or without the addition of cholesterol. Thus, for example, DOPG/DOPC vesicles can be prepared by drying 50 mg each of DOPG and DOPC under a stream of nitrogen gas into a sonication vial. The sample is placed under a vacuum pump overnight and is hydrated the following day with deionized water. The sample is then sonicated for 2 hours in a capped vial, using a Heat Systems model 350 sonicator equipped with an inverted cup (bath type) probe at the maximum setting while the bath is circulated at 15EC. Alternatively, negatively

charged vesicles can be prepared without sonication to produce multilamellar vesicles or by extrusion through nucleopore membranes to produce unilamellar vesicles of discrete size. Other methods are known and available to those of skill in the art.

- The liposomes can comprise multilamellar vesicles (MLVs), small unilamellar [0430] vesicles (SUVs), or large unilamellar vesicles (LUVs), with SUVs being preferred. The various liposome-nucleic acid complexes are prepared using methods well known in the art. See, e.g., Straubinger et al., Methods of Immunology (1983), 101:512-527, which is herein incorporated by reference. For example, MLVs containing nucleic acid can be prepared by depositing a thin film of phospholipid on the walls of a glass tube and subsequently hydrating with a solution of the material to be encapsulated. SUVs are prepared by extended sonication of MLVs to produce a homogeneous population of unilamellar liposomes. The material to be entrapped is added to a suspension of preformed MLVs and then sonicated. When using liposomes containing cationic lipids, the dried lipid film is resuspended in an appropriate solution such as sterile water or an isotonic buffer solution such as 10 mM Tris/NaCl, sonicated, and then the preformed liposomes are mixed directly with the DNA. The liposome and DNA form a very stable complex due to binding of the positively charged liposomes to the cationic DNA. SUVs find use with small nucleic acid fragments. LUVs are prepared by a number of methods, well known in the art. Commonly used methods include Ca2+-EDTA chelation (Papahadjopoulos et al., Biochim. Biophys. Acta (1975) 394:483; Wilson et al., Cell 17:77 (1979); ether injection (Deamer, D. and Bangham, A., Biochim. Biophys. Acta 443:629 (1976); Ostro et al., Biochem. Biophys. Res. Commun. 76:836 (1977); Fraley et al., Proc. Natl. Acad. Sci. USA 76:3348 (1979)); detergent dialysis (Enoch, H. and Strittmatter, P., Proc. Natl. Acad. Sci. USA 76:145 (1979)); and reverse-phase evaporation (REV) (Fraley et al., J. Biol. Chem. 255:10431 (1980); Szoka et al., Proc. Natl. Acad. Sci. USA 75:145 (1978); Schaefer-Ridder et al., Science 215:166 (1982)), which are herein incorporated by reference.
- [0431] Generally, the ratio of DNA to liposomes will be from about 10:1 to about 1:10. Preferably, the ration will be from about 5:1 to about 1:5. More preferably, the ration will be about 3:1 to about 1:3. Still more preferably, the ratio will be about 1:1.
- [0432] U.S. Patent No. 5,676,954 (which is herein incorporated by reference) reports on the injection of genetic material, complexed with cationic liposomes carriers, into mice. U.S.

Patent Nos. 4,897,355, 4,946,787, 5,049,386, 5,459,127, 5,589,466, 5,693,622, 5,580,859, 5,703,055, and international publication no. WO 94/9469 (which are herein incorporated by reference) provide cationic lipids for use in transfecting DNA into cells and mammals. U.S. Patent Nos. 5,589,466, 5,693,622, 5,580,859, 5,703,055, and International Publication No. WO 94/9469 provide methods for delivering DNA-cationic lipid complexes to mammals.

- [0433] In certain embodiments, cells are engineered, ex vivo or *in vivo*, using a retroviral particle containing RNA which comprises a sequence encoding a polypeptide of the present invention. Retroviruses from which the retroviral plasmid vectors may be derived include, but are not limited to, Moloney Murine Leukemia Virus, spleen necrosis virus, Rous sarcoma Virus, Harvey Sarcoma Virus, avian leukosis virus, gibbon ape leukemia virus, human immunodeficiency virus, Myeloproliferative Sarcoma Virus, and mammary tumor virus.
- [0434] The retroviral plasmid vector is employed to transduce packaging cell lines to form producer cell lines. Examples of packaging cells which may be transfected include, but are not limited to, the PE501, PA317, R-2, R-AM, PA12, T19-14X, VT-19-17-H2, RCRE, RCRIP, GP+E-86, GP+envAm12, and DAN cell lines as described in Miller, Human Gene Therapy 1:5-14 (1990), which is incorporated herein by reference in its entirety. The vector may transduce the packaging cells through any means known in the art. Such means include, but are not limited to, electroporation, the use of liposomes, and CaPO₄ precipitation. In one alternative, the retroviral plasmid vector may be encapsulated into a liposome, or coupled to a lipid, and then administered to a host.
- [0435] The producer cell line generates infectious retroviral vector particles which include polynucleotide encoding a polypeptide of the present invention. Such retroviral vector particles then may be employed, to transduce eukaryotic cells, either *in vitro* or *in vivo*. The transduced eukaryotic cells will express a polypeptide of the present invention.
- [0436] In certain other embodiments, cells are engineered, ex vivo or *in vivo*, with polynucleotide contained in an adenovirus vector. Adenovirus can be manipulated such that it encodes and expresses a polypeptide of the present invention, and at the same time is inactivated in terms of its ability to replicate in a normal lytic viral life cycle. Adenovirus expression is achieved without integration of the viral DNA into the host cell chromosome, thereby alleviating concerns about insertional mutagenesis. Furthermore,

adenoviruses have been used as live enteric vaccines for many years with an excellent safety profile (Schwartz, et al., Am. Rev. Respir. Dis.109:233-238 (1974)). Finally, adenovirus mediated gene transfer has been demonstrated in a number of instances including transfer of alpha-1-antitrypsin and CFTR to the lungs of cotton rats (Rosenfeld et al., Science 252:431-434 (1991); Rosenfeld et al., Cell 68:143-155 (1991)). Furthermore, extensive studies to attempt to establish adenovirus as a causative agent in human cancer were uniformly negative (Green et al., Proc. Natl. Acad. Sci. USA 76:6606 (1979)).

- [0437] Suitable adenoviral vectors useful in the present invention are described, for example, in Kozarsky and Wilson, Curr. Opin. Genet. Devel. 3:499-503 (1993); Rosenfeld et al., Cell 68:143-155 (1992); Engelhardt et al., Human Genet. Ther. 4:759-769 (1993); Yang et al., Nature Genet. 7:362-369 (1994); Wilson et al., Nature 365:691-692 (1993); and U.S. Patent No. 5,652,224, which are herein incorporated by reference. For example, the adenovirus vector Ad2 is useful and can be grown in human 293 cells. These cells contain the E1 region of adenovirus and constitutively express Ela and Elb, which complement the defective adenoviruses by providing the products of the genes deleted from the vector. In addition to Ad2, other varieties of adenovirus (e.g., Ad3, Ad5, and Ad7) are also useful in the present invention.
- [0438] Preferably, the adenoviruses used in the present invention are replication deficient. Replication deficient adenoviruses require the aid of a helper virus and/or packaging cell line to form infectious particles. The resulting virus is capable of infecting cells and can express a polynucleotide of interest which is operably linked to a promoter, but cannot replicate in most cells. Replication deficient adenoviruses may be deleted in one or more of all or a portion of the following genes: E1a, E1b, E3, E4, E2a, or L1 through L5.
- [0439] In certain other embodiments, the cells are engineered, ex vivo or *in vivo*, using an adeno-associated virus (AAV). AAVs are naturally occurring defective viruses that require helper viruses to produce infectious particles (Muzyczka, N., Curr. Topics in Microbiol. Immunol. 158:97 (1992)). It is also one of the few viruses that may integrate its DNA into non-dividing cells. Vectors containing as little as 300 base pairs of AAV can be packaged and can integrate, but space for exogenous DNA is limited to about 4.5 kb. Methods for producing and using such AAVs are known in the art. See, for example, U.S.

Patent Nos. 5,139,941, 5,173,414, 5,354,678, 5,436,146, 5,474,935, 5,478,745, and 5,589,377.

- [0440] For example, an appropriate AAV vector for use in the present invention will include all the sequences necessary for DNA replication, encapsidation, and host-cell integration. The polynucleotide construct is inserted into the AAV vector using standard cloning methods, such as those found in Sambrook et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Press (1989). The recombinant AAV vector is then transfected into packaging cells which are infected with a helper virus, using any standard technique, including lipofection, electroporation, calcium phosphate precipitation, etc. Appropriate helper viruses include adenoviruses, cytomegaloviruses, vaccinia viruses, or herpes viruses. Once the packaging cells are transfected and infected, they will produce infectious AAV viral particles which contain the polynucleotide construct. These viral particles are then used to transduce eukaryotic cells, either ex vivo or *in vivo*. The transduced cells will contain the polynucleotide construct integrated into its genome, and will express a polypeptide of the invention.
- [0441] Another method of gene therapy involves operably associating heterologous control regions and endogenous immune/hematopoietic antigen polynucleotide sequences (e.g., encoding an immune/hematopoietic antigen polypeptide of the present invention) via homologous recombination (see, e.g., U.S. Patent No. 5,641,670, issued June 24, 1997; International Publication No. WO 96/29411, published September 26, 1996; International Publication No. WO 94/12650, published August 4, 1994; Koller et al., Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); and Zijlstra et al., Nature 342:435-438 (1989), which are herein incorporated by reference. This method involves the activation of a gene which is present in the target cells, but which is not normally expressed in the cells, or is expressed at a lower level than desired.
- [0442] Polynucleotide constructs are made, using standard techniques known in the art, which contain the promoter with targeting sequences flanking the promoter. Suitable promoters are described herein. The targeting sequence is sufficiently complementary to an endogenous sequence to permit homologous recombination of the promoter-targeting sequence with the endogenous sequence. The targeting sequence will be sufficiently near the 5' end of the desired endogenous polynucleotide sequence so the promoter will be operably linked to the endogenous sequence upon homologous recombination.

[0443] The promoter and the targeting sequences can be amplified using PCR. Preferably, the amplified promoter contains distinct restriction enzyme sites on the 5' and 3' ends. Preferably, the 3' end of the first targeting sequence contains the same restriction enzyme site as the 5' end of the amplified promoter and the 5' end of the second targeting sequence contains the same restriction site as the 3' end of the amplified promoter. The amplified promoter and targeting sequences are digested and ligated together.

- [0444] The promoter-targeting sequence construct is delivered to the cells, either as naked polynucleotide, or in conjunction with transfection-facilitating agents, such as liposomes, viral sequences, viral particles, whole viruses, lipofection, precipitating agents, etc., described in more detail above. The P promoter-targeting sequence can be delivered by any method, included direct needle injection, intravenous injection, topical administration, catheter infusion, particle accelerators, etc. The methods are described in more detail below.
- [0445] The promoter-targeting sequence construct is taken up by cells. Homologous recombination between the construct and the endogenous sequence takes place, such that an endogenous sequence is placed under the control of the promoter. The promoter then drives the expression of the endogenous sequence.
- [0446] The polynucleotide encoding a polypeptide of the present invention may contain a secretory signal sequence that facilitates secretion of the protein. Typically, the signal sequence is positioned in the coding region of the polynucleotide to be expressed towards or at the 5' end of the coding region. The signal sequence may be homologous or heterologous to the immune/hematopoietic antigen polynucleotide of interest and may be homologous or heterologous to the cells to be transfected. Additionally, the signal sequence may be chemically synthesized using methods known in the art.
- [0447] Any mode of administration of any of the above-described polynucleotides constructs can be used so long as the mode results in the expression of one or more molecules in an amount sufficient to provide a therapeutic effect. This includes direct needle injection, systemic injection, catheter infusion, biolistic injectors, particle accelerators (i.e., "gene guns"), gelfoam sponge depots, other commercially available depot materials, osmotic pumps (e.g., Alza minipumps), oral or suppositorial solid (tablet or pill) pharmaceutical formulations, and decanting or topical applications during surgery. For example, direct injection of naked calcium phosphate-precipitated plasmid into rat

liver and rat spleen or a protein-coated plasmid into the portal vein has resulted in gene expression of the foreign gene in the rat livers (Kaneda et al., Science 243:375 (1989)).

- [0448] A preferred method of local administration is by direct injection. Preferably, a recombinant molecule of the present invention complexed with a delivery vehicle is administered by direct injection into or locally within the area of arteries. Administration of a composition locally within the area of arteries refers to injecting the composition centimeters and preferably, millimeters within arteries.
- [0449] Another method of local administration is to contact a polynucleotide construct of the present invention in or around a surgical wound. For example, a patient can undergo surgery and the polynucleotide construct can be coated on the surface of tissue inside the wound or the construct can be injected into areas of tissue inside the wound.
- [0450] Therapeutic compositions useful in systemic administration, include recombinant molecules of the present invention complexed to a targeted delivery vehicle of the present invention. Suitable delivery vehicles for use with systemic administration comprise liposomes comprising ligands for targeting the vehicle to a particular site. In specific embodiments, suitable delivery vehicles for use with systemic administration comprise liposomes comprising polypeptides of the invention for targeting the vehicle to a particular site.
- [0451] Preferred methods of systemic administration, include intravenous injection, aerosol, oral and percutaneous (topical) delivery. Intravenous injections can be performed using methods standard in the art. Aerosol delivery can also be performed using methods standard in the art (see, for example, Stribling et al., Proc. Natl. Acad. Sci. USA 189:11277-11281, 1992, which is incorporated herein by reference). Oral delivery can be performed by complexing a polynucleotide construct of the present invention to a carrier capable of withstanding degradation by digestive enzymes in the gut of an animal. Examples of such carriers, include plastic capsules or tablets, such as those known in the art. Topical delivery can be performed by mixing a polynucleotide construct of the present invention with a lipophilic reagent (e.g., DMSO) that is capable of passing into the skin.
- [0452] Determining an effective amount of substance to be delivered can depend upon a number of factors including, for example, the chemical structure and biological activity of the substance, the age and weight of the animal, the precise condition requiring treatment and its severity, and the route of administration. The frequency of treatments depends

upon a number of factors, such as the amount of polynucleotide constructs administered per dose, as well as the health and history of the subject. The precise amount, number of doses, and timing of doses will be determined by the attending physician or veterinarian.

[0453] Therapeutic compositions of the present invention can be administered to any animal, preferably to mammals and birds. Preferred mammals include humans, dogs, cats, mice, rats, rabbits sheep, cattle, horses and pigs, with humans being particularly preferred.

Biological Activities

- [0454] Polynucleotides or polypeptides, or agonists or antagonists of the present invention, can be used in assays to test for one or more biological activities. If these polynucleotides or polypeptides, or agonists or antagonists of the present invention, do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides and polypeptides, and agonists or antagonists could be used to treat, prevent diagnose and/or prognose the associated disease.
- [0455] The immune/hematopoietic antigen polynucleotides and polypeptides of the invention are predicted to have predominant expression in hematopoietic tissues (e.g., bone marrow, fetal liver, and fetal spleen) or cells and tissues of the immune system (e.g., lymph nodes, spleen, B cells, T cells, monocytes, macrophages, dendritic cells, neutrophils, mast cells, basophils, and eosinophils).
- or cells and tissues of the immune system (e.g., lymph nodes, spleen, B cells, T cells, monocytes, macrophages, dendritic cells, neutrophils, mast cells, basophils, and eosinophils) antigens of the invention may be useful as therapeutic molecules. Each would be useful for diagnosis, detection, treatment and/or prevention of diseases or disorders of the immune system and or diseases and disorders of hematopoiesis, such as anemias, autoimmune diseases, immunodeficiencies, allergies, asthma, and inflammation) and infectious/parasitic diseases.
- [0457] In a preferred embodiment, polynucleotides of the invention (e.g., a nucleic acid sequence of SEQ ID NO:X or the complement thereof; or the cDNA sequence contained in Clone ID NO:Z, or fragments or variants thereof) and/or polypeptides of the invention (e.g., an amino acid sequence contained in SEQ ID NO:Y, an amino acid

sequence encoded by SEQ ID NO:X, or the complement threof, an amino acid sequence encoded by the cDNA sequence contained in Clone ID NO:Z and fragments or variants thereof as described herein) are useful for the diagnosis, detection, treatement, and/or prevention of diseases or disorders of the tissues/cells corresponding to the library source disclosed in column 7 of Table 1A expressing the corresponding immune/hematopoietic sequence disclosed in the same row of Table 1A.

[0458] Particularly, the immune/hematopoietic antigens may be a useful therapeutic for cancers of cells of hematopoitic origin. Treatment, diagnosis, detection, and/or prevention of immune/hematopoietic-related disorders could be carried out using an immune/hematopoietic antigen or soluble form of an immune/hematopoietic antigen, an immune/hematopoietic antigen ligand, gene therapy, or ex vivo applications. Moreover, inhibitors of an immune/hematopoietic antigen, either blocking antibodies or mutant forms, could modulate the expression of the immune/hematopoietic antigen. These inhibitors may be useful to treat, diagnose, detect, and/or prevent diseases associated with the misregulation of an immune/hematopoietic antigen.

In one embodiment, the invention provides a method for the specific delivery of compositions of the invention to cells (e.g., normal or diseased immune/hematopoietic cells) by administering polypeptides of the invention (e.g., immune/hematopoietic antigen polypeptides or anti-immune/hematopoietic antigen antibodies) that are associated with heterologous polypeptides or nucleic acids. In one example, the invention provides a method for delivering a therapeutic protein into the targeted cell (e.g., an aberrant immune/hematopoietic cell or immune/hematopoietic cancer cell). In another example, the invention provides a method for delivering a single stranded nucleic acid (e.g., antisense or ribozymes) or double stranded nucleic acid (e.g., DNA that can integrate into the cell's genome or replicate episomally and that can be transcribed) into the targeted cell.

[0460] In another embodiment, the invention provides a method for the specific destruction of cells (e.g., the destruction of aberrant immune/hematopoietic cells, including, but not limited to, immune/hematopoietic tumor cells) by administering polypeptides of the invention (e.g., immune/hematopoietic antigen polypeptides or fragments thereof, or anti- immune/hematopoietic antigen antibodies) in association with toxins or cytotoxic prodrugs.

[0461] By "toxin" is meant compounds that bind and activate endogenous cytotoxic effector systems, radioisotopes, holotoxins, modified toxins, catalytic subunits of toxins, cytotoxins (cytotoxic agents), or any molecules or enzymes not normally present in or on the surface of a cell that under defined conditions cause the cell's death. Toxins that may be used according to the methods of the invention include, but are not limited to, radioisotopes known in the art, compounds such as, for example, antibodies (or complement fixing containing portions thereof) that bind an inherent or induced endogenous cytotoxic effector system, thymidine kinase, endonuclease, RNAse, alpha toxin, ricin, abrin, Pseudomonas exotoxin A, diphtheria toxin, saporin, momordin, gelonin, pokeweed antiviral protein, alpha-sarcin and cholera toxin. "Toxin" also includes a cytostatic or cytocidal agent, a therapeutic agent or a radioactive metal ion, e.g., alphaemitters such as, for example, ²¹³Bi, or other radioisotopes such as, for example, ¹⁰³Pd, $^{133}\mathrm{Xe},\,^{131}\mathrm{I},\,^{68}\mathrm{Ge},\,^{57}\mathrm{Co},\,^{65}\mathrm{Zn},\,^{85}\mathrm{Sr},\,^{32}\mathrm{P},\,^{35}\mathrm{S},\,^{90}\mathrm{Y},\,^{153}\mathrm{Sm},\,^{153}\mathrm{Gd},\,^{169}\mathrm{Yb},\,^{51}\mathrm{Cr},\,^{54}\mathrm{Mn},\,^{75}\mathrm{Se},\,^{113}\mathrm{Sn},\,^{113}\mathrm{Sm},\,^{11$ ⁹⁰Yttrium, ¹¹⁷Tin, ¹⁸⁶Rhenium, ¹⁶⁶Holmium, and ¹⁸⁸Rhenium; luminescent labels, such as luminol; and fluorescent labels, such as fluorescein and rhodamine, and biotin.

Techniques known in the art may be applied to label antibodies of the [0462] invention. Such techniques include, but are not limited to, the use of bifunctional conjugating agents (see e.g., U.S. Patent Nos. 5,756,065; 5,714,631; 5,696,239; 5,652,361; 5,505,931; 5,489,425; 5,435,990; 5,428,139; 5,342,604; 5,274,119; 4,994,560; and 5,808,003; the contents of each of which are hereby incorporated by reference in its entirety). A cytotoxin or cytotoxic agent includes any agent that is detrimental to cells. Examples include paclitaxol, cytochalasin B, gramicidin D, ethidium bromide, emetine, mitomycin, etoposide, tenoposide, vincristine, vinblastine, colchicin, doxorubicin, daunorubicin, dihydroxy anthracin dione, mitoxantrone, mithramycin, actinomycin D, 1dehydrotestosterone, glucocorticoids, procaine, tetracaine, lidocaine, propranolol, and puromycin and analogs or homologs thereof. Therapeutic agents include, but are not limited to, antimetabolites (e.g., methotrexate, 6-mercaptopurine, 6-thioguanine, cytarabine, 5-fluorouracil decarbazine), alkylating agents (e.g., mechlorethamine, thioepa chlorambucil, melphalan, carmustine (BSNU) and lomustine (CCNU), cyclothosphamide. busulfan, dibromomannitol, streptozotocin, mitomycin C, and cis- dichlorodiamine platinum (II) (DDP) cisplatin), anthracyclines (e.g., daunorubicin (formerly daunomycin) and doxorubicin), antibiotics (e.g., dactinomycin (formerly actinomycin), bleomycin,

mithramycin, and anthramycin (AMC)), and anti-mitotic agents (e.g., vincristine and vinblastine).

[0463] By "cytotoxic prodrug" is meant a non-toxic compound that is converted by an enzyme, normally present in the cell, into a cytotoxic compound. Cytotoxic prodrugs that may be used according to the methods of the invention include, but are not limited to, glutamyl derivatives of benzoic acid mustard alkylating agent, phosphate derivatives of etoposide or mitomycin C, cytosine arabinoside, daunorubisin, and phenoxyacetamide derivatives of doxorubicin.

It will be appreciated that conditions caused by a decrease in the standard or normal level of immune/hematopoietic antigen activity in an individual, particularly in individuals with immune/hematopoietic-related disorders, can be treated by administration of an immune/hematopoietic antigen polypeptide (e.g., such as, for example, the complete immune/hematopoietic antigen polypeptide, the soluble form of the extracellular domain of an immune/hematopoietic antigen polypeptide, or cells expressing the complete protein) or agonist. Thus, the invention also provides a method of treatment of an individual in need of an increased level of immune/hematopoietic antigen activity comprising administering to such an individual a pharmaceutical composition comprising an amount of an isolated immune/hematopoietic antigen polypeptide of the invention, or agonist thereof (e.g., an agonistic anti- immune/hematopoietic antigen antibody), effective to increase the immune/hematopoietic antigen activity level in such an individual.

It will also be appreciated that conditions caused by a increase in the standard or normal level of immune/hematopoietic antigen activity in an individual, particularly in an individual with an immune/hematopoietic-related disorder, can be treated by administration of immune/hematopoietic antigen polypeptides (e.g., such as, for example, the complete immune/hematopoietic antigen polypeptide, the soluble form of the extracellular domain of an immune/hematopoietic antigen polypeptide, or cells expressing the complete protein) or antagonist (e.g., an antagonistic anti-immune/hematopoietic antigen antibody). Thus, the invention also provides a method of treatment of an individual in need of an decreased level of immune/hematopoietic antigen activity comprising administering to such an individual a pharmaceutical composition comprising an amount of an isolated immune/hematopoietic antigen polypeptide of the invention, or antagonist thereof (e.g., an antagonistic anti-immune/hematopoietic antigen antibody),

effective to decrease the immune/hematopoietic antigen activity level in such an individual.

[0466] In certain embodiments, a polypeptide of the invention, or polynucleotides, antibodies, agonists, or antagonists corresponding to that polypeptide, may be used to diagnose and/or prognose diseases and/or disorders associated with the tissue(s) in which the polypeptide of the invention is expressed, including one, two, three, four, five, or more tissues disclosed in Table 1A, column 7 (Tissue Distribution Library Code).

[0467] More generally, polynucleotides, translation products and antibodies corresponding to this gene may be useful for the diagnosis, prognosis, prevention, and/or treatment of diseases and/or disorders associated with the following systems.

Immune Activity

Polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention may be useful in treating, preventing, diagnosing and/or prognosing diseases, disorders, and/or conditions of the immune system, by, for example, activating or inhibiting the proliferation, differentiation, or mobilization (chemotaxis) of immune cells. Immune cells develop through a process called hematopoiesis, producing myeloid (platelets, red blood cells, neutrophils, and macrophages) and lymphoid (B and T lymphocytes) cells from pluripotent stem cells. The etiology of these immune diseases, disorders, and/or conditions may be genetic, somatic, such as cancer and some autoimmune diseases, acquired (e.g., by chemotherapy or toxins), or infectious. Moreover, polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention can be used as a marker or detector of a particular immune system disease or disorder.

[0469] In another embodiment, a polypeptide of the invention, or polynucleotides, antibodies, agonists, or antagonists corresponding to that polypeptide, may be used to treat diseases and disorders of the immune system and/or to inhibit or enhance an immune response generated by cells associated with the tissue(s) in which the polypeptide of the invention is expressed, including one, two, three, four, five, or more tissues disclosed in Table 1A, column 7 (Tissue Distribution Library Code).

Polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the [0470] present invention may be useful in treating, preventing, diagnosing, and/or prognosing immunodeficiencies, including both congenital and acquired immunodeficiencies. Examples of B cell immunodeficiencies in which immunoglobulin levels B cell function and/or B cell numbers are decreased include: X-linked agammaglobulinemia (Bruton's disease), X-linked infantile agammaglobulinemia, X-linked immunodeficiency with hyper IgM, non X-linked immunodeficiency with hyper IgM, X-linked lymphoproliferative syndrome (XLP), agammaglobulinemia including congenital and acquired agammaglobulinemia, adult onset agammaglobulinemia, late-onset agammaglobulinemia, dysgammaglobulinemia, hypogammaglobulinemia, unspecified hypogammaglobulinemia, recessive agammaglobulinemia (Swiss type), Selective IgM deficiency, selective IgA deficiency, selective IgG subclass deficiencies, IgG subclass deficiency (with or without IgA deficiency), Ig deficiency with increased IgM, IgG and IgA deficiency with increased IgM, antibody deficiency with normal or elevated Igs, Ig heavy chain deletions, kappa chain deficiency, B cell lymphoproliferative disorder (BLPD), common variable immunodeficiency (CVID), common variable immunodeficiency (CVI) (acquired), and transient hypogammaglobulinemia of infancy.

[0471] In specific embodiments, ataxia-telangiectasia or conditions associated with ataxia-telangiectasia are treated, prevented, diagnosed, and/or prognosing using the polypeptides or polynucleotides of the invention, and/or agonists or antagonists thereof.

[0472] Examples of congenital immunodeficiencies in which T cell and/or B cell function and/or number is decreased include, but are not limited to: DiGeorge anomaly, severe combined immunodeficiencies (SCID) (including, but not limited to, X-linked SCID, autosomal recessive SCID, adenosine deaminase deficiency, purine nucleoside phosphorylase (PNP) deficiency, Class II MHC deficiency (Bare lymphocyte syndrome), Wiskott-Aldrich syndrome, and ataxia telangiectasia), thymic hypoplasia, third and fourth pharyngeal pouch syndrome, 22q11.2 deletion, chronic mucocutaneous candidiasis, natural killer cell deficiency (NK), idiopathic CD4+ T-lymphocytopenia, immunodeficiency with predominant T cell defect (unspecified), and unspecified immunodeficiency of cell mediated immunity.

[0473] In specific embodiments, DiGeorge anomaly or conditions associated with DiGeorge anomaly are treated, prevented, diagnosed, and/or prognosed using polypeptides or polynucleotides of the invention, or antagonists or agonists thereof.

- [0474] Other immunodeficiencies that may be treated, prevented, diagnosed, and/or prognosed using polypeptides or polynucleotides of the invention, and/or agonists or antagonists thereof, include, but are not limited to, chronic granulomatous disease, Chédiak-Higashi syndrome, myeloperoxidase deficiency, leukocyte glucose-6-phosphate dehydrogenase deficiency, X-linked lymphoproliferative syndrome (XLP), leukocyte adhesion deficiency, complement component deficiencies (including C1, C2, C3, C4, C5, C6, C7, C8 and/or C9 deficiencies), reticular dysgenesis, thymic alymphoplasia-aplasia, immunodeficiency with thymoma, severe congenital leukopenia, dysplasia with immunodeficiency, neonatal neutropenia, short limbed dwarfism, and Nezelof syndromecombined immunodeficiency with Igs.
- [0475] In a preferred embodiment, the immunodeficiencies and/or conditions associated with the immunodeficiencies recited above are treated, prevented, diagnosed and/or prognosed using polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention.
- [0476] In a preferred embodiment polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention could be used as an agent to boost immunoresponsiveness among immunodeficient individuals. In specific embodiments, polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention could be used as an agent to boost immunoresponsiveness among B cell and/or T cell immunodeficient individuals.
- [0477] The polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention may be useful in treating, preventing, diagnosing and/or prognosing autoimmune disorders. Many autoimmune disorders result from inappropriate recognition of self as foreign material by immune cells. This inappropriate recognition results in an immune response leading to the destruction of the host tissue. Therefore, the administration of polynucleotides and polypeptides of the invention that can inhibit an immune response, particularly the proliferation, differentiation, or chemotaxis of T-cells, may be an effective therapy in preventing autoimmune disorders.

[0478] Autoimmune diseases or disorders that may be treated, prevented, diagnosed and/or prognosed by polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention include, but are not limited to, one or more of the following: systemic lupus erythematosus, rheumatoid arthritis, ankylosing spondylitis, multiple sclerosis, autoimmune thyroiditis, Hashimoto's thyroiditis, autoimmune hemolytic anemia, hemolytic anemia, thrombocytopenia, autoimmune thrombocytopenia purpura, autoimmune neonatal thrombocytopenia, idiopathic thrombocytopenia purpura, purpura (e.g., Henloch-Scoenlein purpura), autoimmunocytopenia, Goodpasture's syndrome, Pemphigus vulgaris, myasthenia gravis, Grave's disease (hyperthyroidism), and insulin-resistant diabetes mellitus.

[0479] Additional disorders that are likely to have an autoimmune component that may be treated, prevented, and/or diagnosed with the compositions of the invention include, but are not limited to, type II collagen-induced arthritis, antiphospholipid syndrome, dermatitis, allergic encephalomyelitis, myocarditis, relapsing polychondritis, rheumatic heart disease, neuritis, uveitis ophthalmia, polyendocrinopathies, Reiter's Disease, Stiff-Man Syndrome, autoimmune pulmonary inflammation, autism, Guillain-Barre Syndrome, insulin dependent diabetes mellitus, and autoimmune inflammatory eye disorders.

[0480] Additional disorders that are likely to have an autoimmune component that may be treated, prevented, diagnosed and/or prognosed with the compositions of the invention include, but are not limited to, scleroderma with anti-collagen antibodies (often characterized, e.g., by nucleolar and other nuclear antibodies), mixed connective tissue disease (often characterized, e.g., by antibodies to extractable nuclear antigens (e.g., ribonucleoprotein)), polymyositis (often characterized, e.g., by nonhistone ANA), pernicious anemia (often characterized, e.g., by antiparietal cell, microsomes, and intrinsic factor antibodies), idiopathic Addison's disease (often characterized, e.g., by humoral and cell-mediated adrenal cytotoxicity, infertility (often characterized, e.g., by antispermatozoal antibodies), glomerulonephritis (often characterized, e.g., by glomerular basement membrane antibodies or immune complexes), bullous pemphigoid (often characterized, e.g., by IgG and complement in basement membrane), Sjogren's syndrome (often characterized, e.g., by multiple tissue antibodies, and/or a specific nonhistone ANA (SS-B)), diabetes mellitus (often characterized, e.g., by cell-mediated and humoral islet

cell antibodies), and adrenergic drug resistance (including adrenergic drug resistance with asthma or cystic fibrosis) (often characterized, e.g., by beta-adrenergic receptor antibodies).

Interested, prevented, diagnosed and/or prognosed with the compositions of the invention include, but are not limited to, chronic active hepatitis (often characterized, e.g., by smooth muscle antibodies), primary biliary cirrhosis (often characterized, e.g., by mitochondria antibodies), other endocrine gland failure (often characterized, e.g., by specific tissue antibodies in some cases), vitiligo (often characterized, e.g., by melanocyte antibodies), vasculitis (often characterized, e.g., by Ig and complement in vessel walls and/or low serum complement), post-MI (often characterized, e.g., by myocardial antibodies), cardiotomy syndrome (often characterized, e.g., by myocardial antibodies), urticaria (often characterized, e.g., by IgG and IgM antibodies to IgE), atopic dermatitis (often characterized, e.g., by IgG and IgM antibodies to IgE), asthma (often characterized, e.g., by IgG and IgM antibodies to IgE), argunlomatous, degenerative, and atrophic disorders.

[0482] In a preferred embodiment, the autoimmune diseases and disorders and/or conditions associated with the diseases and disorders recited above are treated, prevented, diagnosed and/or prognosed using for example, antagonists or agonists, polypeptides or polynucleotides, or antibodies of the present invention. In a specific preferred embodiment, rheumatoid arthritis is treated, prevented, and/or diagnosed using polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention.

[0483] In another specific preferred embodiment, systemic lupus erythematosus is treated, prevented, and/or diagnosed using polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention. In another specific preferred embodiment, idiopathic thrombocytopenia purpura is treated, prevented, and/or diagnosed using polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention.

[0484] In another specific preferred embodiment IgA nephropathy is treated, prevented, and/or diagnosed using polynucleotides, polypeptides, antibodies, and/or agonists or antagonists of the present invention.